Sequence Listing

5	<120 <130 <160	N 0 > P P 0 >		ls, 'ied a eptic 29	Timo and : des :	Isola and '	Thera									c Aci	ds and Same
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15	<222 <400	O > 0	(1). 1														
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20	aag Lys	aag Lys	cca Pro	gtt Val 20	cct Pro	gag Glu	att Ile	ctg Leu	cca Pro 25	agg Arg	cca Pro	ccc Pro	cgg Arg	gct Ala 30	ttg Leu	ttc Phe	96
	tgc Cys	ctg Leu	acc Thr 35	ctg Leu	gag Glu	aac Asn	ccc Pro	ctg Leu 40	agg Arg	aag Lys	gcc Ala	tgc Cys	atc Ile 45	agc Ser	att Ile	gta Val	144
25	gaa Glu	tgg Trp 50	aag Lys	ccc Pro	ttc Phe	gag Glu	acg Thr 55	atc Ile	atc Ile	ttg Leu	ctc Leu	acc Thr 60	atc Ile	ttt Phe	gcc Ala	aat Asn	192
30	Cys	gtg Val	gcc Ala	ctg Leu	gcc Ala	gtg Val 70	tac Tyr	ctg Leu	ccc Pro	atg Met	ccg Pro 75	gaa Glu	gat Asp	gac Asp	aac Asn	aac Asn 80	240
	tct Ser	ctg Leu	aac Asn	ctc Leu	ggc Gly 85	ctg Leu	gag Glu	aag Lys	ctg Leu	gag Glu 90	tat Tyr	ttc Phe	ttc Phe	ctc Leu	att Ile 95	gtc Val	288
35	ttc Phe	tcg Ser	att Ile	gaa Glu 100	gcc Ala	gcc Ala	atg Met	aag Lys	atc Ile 105	att Ile	gcc Ala	tac Tyr	ggc Gly	ttc Phe 110	tta Leu	ttc Phe	336
	cac His	cag Gln	gac Asp 115	gct Ala	tac Tyr	ctg Leu	cgc Arg	agt Ser 120	ggc	tgg Trp	aat Asn	gtg Val	ctg Leu 125	gac Asp	ttc Phe	acc Thr	384
40	att Ile	gtc Val 130	ttc Phe	ctg Leu	gly ggg	gtc Val	ttc Phe 135	acc Thr	gtg Val	att Ile	ctg Leu	gaa Glu 140	cag Gln	gtt Val	aac Asn	gtc Val	432
45	atc Ile 145	caa Gln	agc Ser	cac His	aca Thr	gcc Ala 150	cca Pro	atg Met	agc Ser	agc Ser	aaa Lys 155	gga Gly	gcc Ala	ggc Gly	ttg Leu	gat Asp 160	480

	gtc Val	aag Lys	gcc Ala	ctc Leu	aga Arg 165	gcc Ala	ttc Phe	cga Arg	gtg Val	ctc Leu 170	aga Arg	ccc Pro	ctc Leu	cgg Arg	ctg Leu 175	gtg Val	528
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	atg Met	ctc Leu	ccc Pro 195	ctc Leu	ttt Phe	cac His	atc Ile	gcc Ala 200	ctg Leu	ctg Leu	gtc Val	ctc Leu	ttt Phe 205	atg Met	gtc Val	atc Ile	624
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15	acc Thr 225	tgc Cys	tac Tyr	ttc Phe	att Ile	ggt Gly 230	aca Thr	gat Asp	atc Ile	gtg Val	gcc Ala 235	acg Thr	gtg Val	gag Glu	aat Asn	gaa Glu 240	720
	gag Glu	cca Pro	tcg Ser	ccc Pro	tgc Cys 245	gcc Ala	agg Arg	acg Thr	ggc Gly	tca Ser 250	gly ggg	cgc Arg	cgg Arg	tgc Cys	acc Thr 255	atc Ile	768
20	aat Asn	ggc Gly	agt Ser	gag Glu 260	tgc Cys	cgg Arg	ggc Gly	ggc	tgc Cys 265	cca Pro	gly ggg	ccc Pro	aac Asn	cat His 270	ggc	atc Ile	816
	acc Thr	cac His	ttc Phe 275	gac Asp	aac Asn	ttc Phe	ggc Gly	ttc Phe 280	tcc Ser	atg Met	ctc Leu	acc Thr	gtg Val 285	tac Tyr	cag Gln	tgc Cys	864
25	att Ile	acc Thr 290	atg Met	gag Glu	gga Gly	tgg Trp	act Thr 295	gac Asp	gtc Val	ctt Leu	tac Tyr	tgg Trp 300	gtc Val	aat Asn	gat Asp	gcc Ala	912
30	atc Ile 305	gly aaa	aat Asn	gag Glu	tgg Trp	ccc Pro 310	tgg Trp	atc Ile	tat Tyr	ttt Phe	gtc Val 315	acc Thr	ctc Leu	att Ile	ttg Leu	ctg Leu 320	960
	gga Gly	tcc Ser	ttc Phe	ttc Phe	atc Ile 325	ctc Leu	aac Asn	ctg Leu	gtg Val	ctg Leu 330	ggt Gly	gtc Val	ctg Leu	agt Ser	999 335	gaa Glu	1008
35	ttc Phe	acc Thr	aag Lys	gag Glu 340	cgg Arg	gag Glu	aag Lys	gcc Ala	aag Lys 345	tcc Ser	agg Arg	gga Gly	acc Thr	ttc Phe 350	cag Gln	aag Lys	1056
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45	Lys	ctg Leu	tct Ser	ttg Leu	Asp	gaa Glu 390	ggt Gly	ggc	tct Ser	Asp	aca Thr 395	gag Glu	agc Ser	ctg Leu	tat Tyr	gaa Glu 400	1200

		-		_		aaa Lys			_			_				_	1248
5						cgc Arg											1296
						gtg Val											1344
10						cac His											1392
15						gtg Val 470											1440
						ctg Leu											1488
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						ctc Leu											1728
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15	ctg Leu 705	gag Glu	cag Gln	aaa Lys	ccc Pro	aag Lys 710	ggt Gly	gag Glu	ggc Gly	atc Ile	ccc Pro 715	acc Thr	act Thr	gcc Ala	aag Lys	ctg Leu 720	2160
	aaa Lys	atc Ile	gat Asp	gag Glu	ttt Phe 725	gaa Glu	tct Ser	aat Asn	gtc Val	aat Asn 730	gag Glu	gtg Val	aag Lys	gat Asp	ccc Pro 735	tac Tyr	2208
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20	gac Asp	Gl ^A aaa	gac Asp	ccc Pro 980	atg Met	cag Gln	ata Ile	gag Glu	ctg Leu 985	cgt Arg	cac His	cgc Arg	gag Glu	tgg Trp 990	gta Val	cac His	2976
	agc Ser	gac Asp	ttc Phe 995	cac His	ttc Phe	gac Asp	aat Asn	gtg Val 100	Le	c to u Se	a go r Al	c at .a Me	et Me	g t t S 05		tc ttc eu Phe	3024
25	acg Thr	_	tc			c ga		a t	.gg c	ct c	30 0			+	220		
		Val		L 111.		0 01		у Т 15	rp P	ro G	ln I	eu I	etg Leu L020		Lys		3069
30	Ile		0 tc Se	c aa	t gc	g ga	10 g ga u As	15 c g	rp P tg g al G	ro G	ln I	leu I itc t ile T	Leu L020 Cac	Tyr aac		Ala cgt	3114
30	Ile) qtq	gac Asp 102	tc Se 5 At	c aa r As: g gc	t gc n Al	g ga a Gl	g ga u As 10 c tt e Ph	15 c g p V 30	rtg g Mal G	gt cally F	de la company de	atc tile i	Leu L020 Lac Tyr	Tyr aac Asn	Lys aac Asn	Ala cgt Arg gcc	
	Ile gtg Val	gac Asp 102 gag Glu 104	0 tc Se 5 at Me 0 at	c aa r As: g gc t Al	t gc n Al c at a Il	g ga a Gl c tt e Ph c at	g gau As 10 c tte Ph c tte Ph	15 C 9 P V 30 C a e I 45	rtg gral G	gt of the state of	In I	atc tatc atc atc atc atc atc atc atc atc	Leu 1020 cac Tyr 1035 atc	Tyr aac Asn ctc Leu	aac Asn att Ile	ala cgt Arg gcc Ala ttc	3114
	gtg Val ttc Phe	gac Asp 102 gag Glu 104 ttc Phe 105 gag	tc se 5 at Me 0 at Me 5	c aa r As g gc t Al g at t Me	t gc n Al c at a Il g aa t As	g ga a Gl c tt e Ph c at n Il	g ga u As 10 c tt e Ph 10 c tt e Ph 10 t ga r Gl	15 c g p V 30 c a 45 t g 60	rtg gral G	gt of the state of	in I	atc tile i	Leu LO20 Cac Fyr LO35 atc Ile LO50 att Ile	aac Asn ctc Leu gtc Val	aac Asn att Ile acc Thr	cgt Arg gcc Ala ttc Phe	3114 3159
35	gtg Val ttc Phe cag Gln	gac Asp 102 gag Glu 104 ttc Phe 105 gag Glu 107 cag	tc se se at Me o at Me ca Gl	c aac r Asc g gc t Al g at t Me g gg n Gl	t gc n Al c at a Il g aa t As a ga y Gl a tg	g gaa Glic tte Phocatn Il gacu Th	g ga u As 10 c tt e Ph 10 c tt e Ph 10 t ga r Gl a ca l Gl	15 c p V 30 c e I 45 t e V 175	tg g fal G ftg g fal G f	gt of the state of	cc ac a	atc to the state atc as the state at a state	Leu LO20 Cac Tyr LO35 atc Ile LO50 att Ile 1065	Tyr aac Asn ctc Leu gtc Val ctg Leu cgc	aac Asn att Ile acc Thr	cgt Arg gcc Ala ttc Phe aag Lys	3114 3159 3204

	att Ile	gtc Val 1115	Thr	tcc Ser	tcc Ser	tac Tyr	ttt Phe 1120	Glu	tac Tyr	ctg Leu	atg Met	ttt Phe 1125	Ala	ctc Leu	atc Ile		3384
5		ctc Leu 1130	Asn	acc Thr	atc	tgc Cys	ctc Leu 1135	Gly	atg Met	cag Gln	cac His	tac Tyr 1140	Asn	cag Gln	tcg Ser		3429
	gag Glu	cag Gln 1145	Met	aac Asn	cac His	atc Ile	tca Ser 1150	Asp	atc Ile	ctc Leu	aat Asn	gtg Val 1155	Ala	ttc Phe	act Thr		3474
10	atc Ile	atc Ile 1160	Phe	acc Thr	ctg Leu	gag Glu	atg Met 1165	Ile	ctc Leu	aag Lys	ctc Leu	atg Met 1170	Ala	ttc Phe	aag Lys		3519
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	att Ile	gtc Val 1190	Ile	ggc	agc Ser	atc Ile	att Ile 1195	Asp	gtc Val	atc Ile	ctc Leu	agt Ser 1200	gag Glu	atc Ile	gac Asp		3609
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30	Pro	tac Tyr 1250	gtg Val	gct Ala	ctg Leu	ctc Leu	atc Ile 1255	gtc Val	atg Met	ctc Leu	ttc Phe	ttc Phe 1260	atc Ile	tac Tyr	_		3789
	gtc Val	atc Ile 1265	ggc Gly	atg Met	cag Gln	atg Met	ttt Phe 1270	Gly 999	aag Lys	atc Ile	gcc Ala	ttg Leu 1275	gtg Val	gat Asp	gly ggg		3834
35	acc Thr	caa Gln 1280	ata Ile	aac Asn	cgg Arg	aac Asn	aac Asn 1285	aac Asn	ttc Phe	cag Gln	acc Thr	ttc Phe 1290	cca Pro	caa Gln	gct Ala	:	3879
	gtg Val	cta Leu 1295	ctg Leu	ctc Leu	ttc Phe	agg Arg	tgt Cys 1300	gca Ala	aca Thr	ggt Gly	gag Glu	gcc Ala 1305	tgg Trp			:	3924
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			Asp					Gly				ctg Leu 1380		gag Glu		4149
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15	aaa Lys	cac His 1400	ctg Leu	gac Asp	gtg Val	gtg Val	acc Thr 1405	ctg Leu	ctg Leu	aga Arg	agg Arg	att Ile 1410	cag Gln	ccc Pro		4239
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	gaa Glu	agg Arg 1595	Thr	aac Asn	tcc Ser	ctg Leu	ccc Pro 1600	Pro	gtc Val	atg Met	gcc Ala	aat Asn 1605	cag Gln	aga Arg	ccc Pro	4824
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15	ttc Phe	ttg Leu 1625	gag Glu	gac Asp	ttc Phe	cca Pro	caa Gln 1630	Asp	cca Pro	cgc Arg	acc Thr	aac Asn 1635	ccc Pro	ctg Leu	gct Ala	4914
	cgt Arg	gcc Ala 1640	aat Asn	acc Thr	aac Asn	aat Asn	gcc Ala 1645	aac Asn	gcc Ala	aat Asn	gtc Val	gcc Ala 1650	tat Tyr		aac Asn	4959
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Met Leu Pro Leu Phe His Ile Ala Leu Leu Val Leu Phe Met Val Ile 195 200 205

Ile Tyr Ala Ile Ile Gly Leu Glu Leu Phe Lys Gly Lys Met His Lys 10 210 215 220

Thr Cys Tyr Phe Ile Gly Thr Asp Ile Val Ala Thr Val Glu Asn Glu 225 230 235 240

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Gly Ser Phe Phe Ile Leu Asn Leu Val Leu Gly Val Leu Ser Gly Glu 325 330 335

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Trp Ile Thr Gln Gly Glu Val Met Asp Val Glu Asp Phe Arg Glu Gly 30 370 375 380

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5 Trp Asn Arg Ile Phe Arg Trp Lys Cys His Asp Ile Val Lys Ser Lys
420 425 430

Val Phe Tyr Trp Leu Val Ile Leu Ile Val Ala Leu Asn Thr Leu Ser 435 440 445

Ile Ala Ser Glu His His Asn Gln Pro His Trp Leu Thr Arg Leu Gln 450 455 460

Asp Ile Ala Asn Arg Val Leu Leu Ser Leu Phe Thr Thr Glu Met Leu 465 470 475 480

Met Lys Met Tyr Gly Leu Gly Leu Arg Gln Tyr Phe Met Ser Ile Phe 485 490 495

15 Asn Arg Phe Asp Cys Phe Val Val Cys Ser Gly Ile Leu Glu Ile Leu 500 510

Leu Val Glu Ser Gly Ala Met Thr Pro Leu Gly Ile Ser Val Leu Arg 515 520 525

Cys Ile Arg Leu Leu Arg Ile Phe Lys Ile Thr Lys Tyr Trp Thr Ser 20 535 540

Leu Ser Asn Leu Val Ala Ser Leu Leu Asn Ser Ile Arg Ser Ile Ala 545 550 560

Ser Leu Leu Leu Leu Phe Leu Phe Ile Val Ile Phe Arg Leu Leu 565 570 575

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Arg Arg Ser Asn Phe Asp Asn Phe Pro Gln Ala Leu Ile Ser Val Phe 595 600 605

Gln Val Leu Thr Gly Glu Asp Trp Thr Ser Met Met Tyr Asn Gly Ile 30 $\,$ 610 $\,$ 620

Met Ala Ser Ser Gly Pro Ser Tyr Pro Gly Met Leu Val Cys Ile Tyr 625 630 635 640

Phe Ile Ile Leu Phe Val Cys Gly Asn Tyr Ile Leu Leu Asn Val Phe 645 650 655

5 Leu Ala Ile Ala Val Asp Asn Leu Ala Glu Ala Glu Ser Leu Thr Ser 660 665 670

Ala Gln Lys Ala Lys Ala Glu Glu Lys Lys Arg Arg Lys Met Ser Lys 675 680 685

Gly Leu Pro Asp Lys Ser Glu Glu Glu Lys Ser Thr Met Ala Lys Lys 10 690 695 700

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- Arg Cys Ser Ala Pro Ala Thr Ala Leu Leu Ile Gln Lys Ala Leu 30 1760 1765 1770

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55 60

Ala Ala Gln Thr Met Ser Thr Ser Ala Pro Pro Pro Val Gly Ser Leu 65 70 75 80

Ser Gln Arg Lys Arg Gln Gln Tyr Ala Lys Ser Lys Lys Gln Gly Asn 85 90 95

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Asn Pro Ile Arg Arg Ala Cys Ile Ser Ile Val Glu Trp Lys Pro Phe 115 120 125

Asp Ile Phe Ile Leu Leu Ala Ile Phe Ala Asn Cys Val Ala Leu Ala 20 130 135 140

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 - Phe Ser Pro Thr Asn Lys Ile Arg Val Leu Cys His Arg Ile Val Asn 100 105 110
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 - Ser Phe Cys Arg Asn Tyr Phe Asn Met Leu Asp Leu Leu Val Val Ala 180 185 190
- Val Ser Leu Ile Ser Met Gly Leu Glu Ser Ser Ala Ile Ser Val Val 20 195 200 205
 - Lys Ile Leu Arg Val Leu Arg Val Leu Arg Pro Leu Arg Ala Ile Asn 210 $\,$ 215 $\,$ 220 $\,$
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 - Met Met Ser Leu Phe Thr Val Ser Thr Phe Glu Gly Trp Pro Gln Leu 370 375 380
 - Leu Tyr Lys Ala Ile Asp Ser Asn Ala Glu Asp Val Gly Pro Ile Tyr 385 390 395 400
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- Phe Gln Glu Gln Gly Glu Thr Glu Tyr Lys Asn Cys Glu Leu Asp Lys 20 435 440 445
 - Asn Gln Arg Gln Cys Val Gln Tyr Ala Leu Lys Ala Arg Pro Leu Arg 450 455 460
 - Cys Tyr Ile Pro Lys Asn Pro Tyr Gln Tyr Gln Val Trp Tyr Ile Val 465 470 475 480
- 25 Thr Ser Ser Tyr Phe Glu Tyr Leu Met Phe Ala Leu Ile Met Leu Asn 485 490 495
 - Thr Ile Cys Leu Gly Met Gln His Tyr Asn Gln Ser Glu Gln Met Asn 500 505 510
- His Ile Ser Asp Ile Leu Asn Val Ala Phe Thr Ile Ile Phe Thr Leu 30 515 520 525

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Leu Tyr Cys Leu Gly Gly Cys Gly Asn Val Asp Pro Asp Glu Ser 580 585 590

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Thr Phe Pro Gln Ala Val Leu Leu Leu Phe Arg Cys Ala Thr Gly Glu 675 680 685

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Val Ile Asn Leu Phe Val Ala Val Ile Met Asp Asn Phe Asp Tyr Leu 740 750

Thr Arg Asp Trp Ser Ile Leu Gly Pro His His Leu Asp Glu Phe Lys 30 755 760 765

- Ala Ile Trp Ala Glu Tyr Asp Pro Glu Ala Lys Gly Arg Ile Lys His
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- 5 Gly Lys Phe Cys Pro His Arg Val Ala Cys Lys Arg Leu Val Gly Met 805 810 815
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 - Glu Gln Ala Asn Glu Glu Leu Arg Ala Ile Ile Lys Lys Ile Trp Lys 850 855 860
 - Arg Thr Ser Met Lys Leu Leu Asp Gln Val Ile Pro Pro Ile Gly Asp 865 870 875 886
- 15 Asp Glu Val Thr Val Gly Lys Phe Tyr Ala Thr Phe Leu Ile Gln Glu 885 890 895
 - His Phe Arg Lys Phe Met Lys Arg Gln Glu Glu Tyr Tyr Gly Tyr Arg 900 905 910
- Pro Lys Lys Asp Ile Val Gln Ile Gln Ala Gly Leu Arg Thr Ile Glu 20 915 920 925
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- $25\,\mathrm{Gly}$ Ile Phe Arg Arg Thr Gly Gly Leu Phe Gly Gln Val Asp Asn Phe 965 970 975
 - Leu Glu Arg Thr Asn Ser Leu Pro Pro Val Met Ala Asn Gln Arg Pro 980 985
- Leu Gln Phe Ala Glu Ile Glu Met Glu Glu Met Glu Ser Pro Val Phe $30 \hspace{1.5cm} 995 \hspace{1.5cm} 1000 \hspace{1.5cm} 1005$

Leu Glu Asp Phe Pro Gln Asp Pro Arg Thr Asn Pro Leu Ala Arg 1015 Ala Asn Thr Asn Asn Ala Asn Ala Asn Val Ala Tyr Gly Asn Ser 1025 1030 1035 ${f 5}$ Asn His Ser Asn Ser His Val $\,$ Phe Ser Ser Val His $\,$ Tyr Glu Arg 1045 1050 1040 Glu Phe Pro Glu Glu Thr Glu Thr Pro Ala Thr Arg Gly Arg Ala 1055 1060 Leu Gly Gln Pro Cys Arg Val Xaa Xaa Xaa Xaa Xaa Xaa Xaa 10 1070 1075 1080 1085 1090 1095 1100 1105 1110 1125 1115 1120 1130 1135 1140 20 1145 1150 1160 Xaa Xaa Thr Gly Pro His Ser Lys Pro Cys Val Glu Met Leu Lys 1175 1180 25 Gly Leu Leu Thr Gln Arg Ala Met Pro Arg Gly Gln Ala Pro Pro 1190 1195 1200 Ala Pro Cys Gln Cys Pro Arg Val Glu Ser Ser Met Pro Glu Asp 1205 1210 1215 Arg Lys Ser Ser Thr Pro Gly Ser Leu His Glu Glu Thr Pro His 30 1220 1225 1230

	Ser	Arg 1235		Thr	Arg	Glu	Asn 1240		s Ser	Arg	, Cys	Ser 1245	Ala	Pro	Ala	
	Thr	Ala 1250		Leu	Ile	Gln	Lys 1255		a Leu	val	Arg	Gly 1260	Gly	Leu	Gly	
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	Leu	Ala 1280		Ala	Суз	Gln	Met 1285		ı Pro	Glu	ı Glu	Val 1290	Glu	Ile	Met	
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	Ser	Ser 1310		Gly	Cys	Leu	Asn 1315		ı Gly	Ser	Ser	Leu 1320	Gly	Ser	Leu	
	Asp	Gln 1325		Gln	Gly	Ser	Gln 1330		ı Thr	Leu	ı Ile	Pro 1335	Pro	Arg	Leu	
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				-	Glu		_		_	ys C	_	cc aag		_	_	285
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		ctg Leu															525
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				atc Ile	_			_			_					1485
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				cag Gln	_			_	_	_		_		_		1629
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	_						_			_		_		_	cag Gln		1965
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		_	_				_	_					_	_	tcg Ser	_	2109
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	-		_			_		_	_	_	_	_	_		cag Gln		2349
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Glu Trp Lys Pro Phe Glu Thr Ile Ile Leu Leu Thr Ile Phe Ala Asn 50 55 60

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Ser Leu Asn Leu Gly Leu Glu Lys Leu Glu Tyr Phe Phe Leu Ile Val 85 90 95

Phe Ser Ile Glu Ala Ala Met Lys Ile Ile Ala Tyr Gly Phe Leu Phe 100 100 105 110

His Gln Asp Ala Tyr Leu Arg Ser Gly Trp Asn Val Leu Asp Phe Thr
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Val Lys Ala Leu Arg Ala Phe Arg Val Leu Arg Pro Leu Arg Leu Val

Ser Gly Val Pro Ser Leu Gln Val Val Leu Asn Ser Ile Phe Lys Ala 180 180 185 190

Met Leu Pro Leu Phe His Ile Ala Leu Leu Val Leu Phe Met Val Ile 195 200 205 Ile Tyr Ala Ile Ile Gly Leu Glu Leu Phe Lys Gly Lys Met His Lys 210 215 220

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Trp Asn Arg Ile Phe Arg Trp Lys Cys His Asp Ile Val Lys Ser Lys 420 425 430

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- 5 Met Lys Met Tyr Gly Leu Gly Leu Arg Gln Tyr Phe Met Ser Ile Phe 485 490 495
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- Leu Val Glu Ser Gly Ala Met Thr Pro Leu Gly Ile Ser Val Leu Arg
 515 520 525
 - Cys Ile Arg Leu Leu Arg Ile Phe Lys Ile Thr Lys Tyr Trp Thr Ser 530 540
 - Leu Ser Asn Leu Val Ala Ser Leu Leu Asn Ser Ile Arg Ser Ile Ala 545 550 555 560
- 15 Ser Leu Leu Leu Leu Phe Leu Phe Ile Val Ile Phe Arg Leu Leu 565 570 575
 - Gly Met Gln Leu Phe Gly Gly Arg Tyr Asp Phe Glu Asp Thr Glu Val 580 585 590
- Arg Arg Ser Asn Phe Asp Asn Phe Pro Gln Ala Leu Ile Ser Val Phe 20 595 600 605
 - Gln Val Leu Thr Gly Glu Asp Trp Thr Ser Met Met Tyr Asn Gly Ile 610 615 620
 - Met Ala Ser Ser Gly Pro Ser Tyr Pro Gly Met Leu Val Cys Ile Tyr 625 630 635 640
- 25 Phe Ile Ile Leu Phe Val Cys Gly Asn Tyr Ile Leu Leu Asn Val Phe 645 650 655
 - Leu Ala Ile Ala Val Asp Asn Leu Ala Glu Ala Glu Ser Leu Thr Ser 660 665 670
- Ala Gln Lys Ala Lys Ala Glu Glu Lys Lys Arg Arg Lys Met Ser Lys 675 680 685

- Gly Leu Pro Asp Lys Ser Glu Glu Glu Lys Ser Thr Met Ala Lys Lys 690 695 700
- Leu Glu Gln Lys Pro Lys Gly Glu Gly Ile Pro Thr Thr Ala Lys Leu 705 710 715 720
- 5 Lys Ile Asp Glu Phe Glu Ser Asn Val Asn Glu Val Lys Asp Pro Tyr
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 - Pro Ser Ala Asp Phe Pro Gly Asp Asp Glu Glu Asp Glu Pro Glu Ile 740 745 750
- Pro Leu Ser Pro Arg Pro Arg Pro Leu Ala Glu Leu Gln Leu Lys Glu
 755 760 765
 - Lys Ala Val Pro Ile Pro Glu Ala Ser Ser Phe Phe Ile Phe Ser Pro 770 775 780
 - Thr Asn Lys Ile Arg Val Leu Cys His Arg Ile Val Asn Ala Thr Trp 785 790 795 800
- 15 Phe Thr Asn Phe Ile Leu Leu Phe Ile Leu Leu Ser Ser Ala Ala Leu 805 810 815
 - Ala Ala Glu Asp Pro Ile Arg Ala Asp Ser Met Arg Asn Gln Ile Leu 820 825 830
- Lys His Phe Asp Ile Gly Phe Thr Ser Val Phe Thr Val Glu Ile Val
 - Leu Lys Met Thr Thr Tyr Gly Ala Phe Leu His Lys Gly Ser Phe Cys 850 855 860
 - Arg Asn Tyr Phe Asn Met Leu Asp Leu Leu Val Val Ala Val Ser Leu 865 870 875 880
- 25 Ile Ser Met Gly Leu Glu Ser Ser Ala Ile Ser Val Val Lys Ile Leu 885 890 895
 - Arg Val Leu Arg Val Leu Arg Pro Leu Arg Ala Ile Asn Arg Ala Lys 900 905 910
- Gly Leu Lys His Val Ala Arg Cys Met Phe Val Ala Ile Ser Thr Ile 30 915 920 925

- Gly Asn Ile Val Leu Val Thr Thr Leu Leu Gln Phe Met Phe Ala Cys 930 935 940
- Ile Gly Val Gln Leu Phe Lys Gly Lys Phe Phe Arg Cys Thr Asp Leu 945 950 955 960
- 5 Ser Lys Met Thr Glu Glu Glu Cys Arg Gly Tyr Tyr Tyr Val Tyr Lys
 965 970 975
 - Asp Gly Asp Pro Met Gln Ile Glu Leu Arg His Arg Glu Trp Val His 980 985 990
- Ser Asp Phe His Phe Asp Asn Val Leu Ser Ala Met Met Ser Leu Phe 10 995 1000 1005
 - Thr Val Ser Thr Phe Glu Gly Trp Pro Gln Leu Leu Tyr Lys Ala 1010 1015 1020
 - Ile Asp Ser Asn Ala Glu Asp Val Gly Pro Ile Tyr Asn Asn Arg 1025 1030 1035
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 - Phe Phe Met Met Asn Ile Phe Val Gly Phe Val Ile Val Thr Phe 1055 1060 1065
- Gln Glu Gln Gly Glu Thr Glu Tyr Lys Asn Cys Glu Leu Asp Lys 20 1070 1080
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 - Arg Cys Tyr Ile Pro Lys Asn Pro Tyr Gln Tyr Gln Val Trp Tyr 1100 1105 1110
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- Ile Ile Phe Thr Leu Glu Met Ile Leu Lys Leu Met Ala Phe Lys 1160 1165 1170
- Ala Arg Gly Tyr Phe Gly Asn Pro Trp Asn Val Phe Asp Phe Leu 1175 1180 1185
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- Cys Gly Asn Val Asp Pro Asp Glu Ser Ala Arg Ile Ser Ser Ala 10 1220 1225 1230
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- Arg Ala Glu Gly Val Arg Thr Leu Leu Trp Thr Phe Ile Lys Ser 1250 1255 1260
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1600

Phe Asp Tyr Leu Thr Arg Asp Trp Ser Ile Leu Gly Pro His His 1385 1390 1395 Leu Asp Glu Phe Lys Ala Ile Trp Ala Glu Tyr Asp Pro Glu Ala 1405 1410 5 Lys Gly Arg Ile Lys His Leu Asp Val Val Thr Leu Leu Arg Arg 1415 1420 Ile Gln Pro Pro Leu Gly Phe Gly Lys Phe Cys Pro His Arg Val 1430 1435 Ala Cys Lys Arg Leu Val Gly Met Asn Met Pro Leu Asn Ser Asp 10 1445 1450 Gly Thr Val Thr Phe Asn Ala Thr Leu Phe Ala Leu Val Arg Thr 1460 1465 Ala Leu Lys Ile Lys Thr Glu Gly Asn Phe Glu Gln Ala Asn Glu 1475 1480 1485 15 Glu Leu Arg Ala Ile Ile Lys Lys Ile Trp Lys Arg Thr Ser Met 1490 1495 1500 Lys Leu Leu Asp Gln Val Ile Pro Pro Ile Gly Asp Asp Glu Val 1505 1510 1515 Thr Val Gly Lys Phe Tyr Ala Thr Phe Leu Ile Gln Glu His Phe 20 1520 1525 Arg Lys Phe Met Lys Arg Gln Glu Glu Tyr Tyr Gly Tyr Arg Pro 1535 1540 Lys Lys Asp Ile Val Gln Ile Gln Ala Gly Leu Arg Thr Ile Glu 1550 $25\,\mathrm{Glu}$ Glu Ala Ala Pro Glu Ile Cys Arg Thr Val Ser Gly Asp Leu 1565 1570 Ala Ala Glu Glu Leu Glu Arg Ala Met Val Glu Ala Ala Met 1580 1585 Glu Glu Gly Ile Phe Arg Arg Thr Gly Gly Leu Phe Gly Gln Val

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Asp Asn Phe Leu Glu Arg Thr Asn Ser Leu Pro Pro Val Met Ala 1610 1615 1620 Asn Gln Arg Pro Leu Gln Phe Ala Glu Ile Glu Met Glu Glu Met 1630 1635 5 Glu Ser Pro Val Phe Leu Glu Asp Phe Pro Gln Asp Pro Arg Thr 1645 Asn Pro Leu Ala Arg Ala Asn Thr Asn Asn Ala Asn Ala Asn Val 1655 1660 Ala Tyr Ala Asn Ser Asn His Ser Asn Ser His Val Phe Ser Ser 10 1670 1675 Val His Tyr Glu Arg Glu Phe Pro Glu Glu Thr Glu Thr Pro Ala 1685 1690 Thr Arg Gly Arg Ala Leu Gly Gln Pro Cys Arg Ser Leu Gly Pro 1700 1705 1710 15 His Ser Lys Pro Cys Val Glu Met Leu Lys Gly Leu Leu Thr Gln 1715 1720 Arg Ala Met Pro Arg Gly Gln Ala Pro Pro Ala Pro Cys Gln Cys 1730 1740 Pro Arg Val Glu Ser Ser Met Pro Glu Asp Arg Lys Ser Ser Thr 20 1745 1750 1755 Pro Gly Ser Leu His Glu Glu Thr Pro His Ser Arg Ser Thr Arg 1765 1770 1760 Glu Asn Thr Ser Arg Cys Ser Ala Pro Ala Thr Ala Leu Leu Ile 1775 1780 25 Gln Lys Ala Leu Val Arg Gly Gly Leu Gly Thr Leu Ala Ala Asp 1790 1795 Ala Asn Phe Ile Met Ala Thr Gly Gln Ala Leu Gly Asp Ala Cys 1805 1810 1815 Gln Met Glu Pro Glu Glu Val Glu Ile Met Ala Thr Glu Leu Leu

1825

Lys Gly Arg Glu Ala Pro Asp Gly Met Ala Ser Ser Leu Gly Cys 1835 1840 Leu Asn Leu Gly Ser Ser Leu Gly Ser Leu Asp Gln His Gln Gly 1850 1855 1860 5 Ser Gln Glu Thr Leu Ile Pro Pro Arg Leu 1865 1870 <210> 13 <211> 6160 <212> DNA 10 <213> Homo sapiens <220> <221> gene <222> (1)..(6160) <220> 15 <221> CDS <222> (226)..(5847) <400> 13 tcaggccggc agcggggagc cgagtggagg ctaattttac ttgctgggag cgaggagagt 60 aatceteetg cececacte etgeceegee eeetggetgg etcageaggg cageteagee 20 gacageetea gecageetag teeceaagge gggggeattg gggacaeagg gaagggaaag 180 cactggggtg ggggagcagg agaaagccag attcccaggg aagcc atg gag cca tcc 237 Met Glu Pro Ser 1 tca ccc cag gat gaa ggc ctg agg aag aaa cag ccc aag aag cca gtt 285 25 Ser Pro Gln Asp Glu Gly Leu Arg Lys Lys Gln Pro Lys Lys Pro Val 5 cct gag att ctg cca agg cca ccc cgg gct ttg ttc tgc ctg acc ctg 333 Pro Glu Ile Leu Pro Arg Pro Pro Arg Ala Leu Phe Cys Leu Thr Leu $30\;\mathrm{gag}$ aac ccc ctg agg aag gcc tgc atc agc att gta gaa tgg aag ccc 381 Glu Asn Pro Leu Arg Lys Ala Cys Ile Ser Ile Val Glu Trp Lys Pro 40 ttc gag acg atc atc ttg ctc acc atc ttt gcc aat tgt gtg gcc ctg 429 Phe Glu Thr Ile Ile Leu Leu Thr Ile Phe Ala Asn Cys Val Ala Leu 35 55 gcc gtg tac ctg ccc atg ccg gaa gat gac aac aac tct ctg aac ctc 477 Ala Val Tyr Leu Pro Met Pro Glu Asp Asp Asn Asn Ser Leu Asn Leu 70 ggc ctg gag aag ctg gag tat ttc ttc ctc att gtc ttc tcg att gaa 525 40 Gly Leu Glu Lys Leu Glu Tyr Phe Phe Leu Ile Val Phe Ser Ile Glu 85 gcc gcc atg aag atc att gcc tac ggc ttc tta ttc cac cag gac gct 573 Ala Ala Met Lys Ile Ile Ala Tyr Gly Phe Leu Phe His Gln Asp Ala

110

	tac Tyr	cto Lev	g cgc ı Arg	agt Ser 120	. Gly	tgg Trp	aat Asn	gtg Val	ctg Leu 125	Asp	tto Phe	acc Thr	att	gto Val	. Phe	ctg Leu	621
Ę	939 939	gto Val	tto Phe	Thr	gtg Val	att Ile	ctg Leu	gaa Glu 140	Gln	gtt Val	aac Asn	gto Val	ato Ile 145	Gln	ago Ser	cac His	669
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	gga Gly	gag Glu	act Thr	gag Glu 1075	Tyr	aag Lys	aac Asn	tgt Cys	gag Glu 1080	Leu	gac Asp	aag Lys	aac Asn	cag Gln 1085	Arg	3483
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Met Glu Pro Ser Ser Pro Gln Asp Glu Gly Leu Arg Lys Lys Gln Pro 5

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 - Val Lys Ala Leu Arg Ala Phe Arg Val Leu Arg Pro Leu Arg Leu Val 165 170 175
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 - Met Leu Pro Leu Phe His Ile Ala Leu Leu Val Leu Phe Met Val Ile 195 200 205
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 - Gly Ser Phe Phe Ile Leu Asn Leu Val Leu Gly Val Leu Ser Gly Glu 325 330 335
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 - Ile Ala Gly Leu Asn Lys Ile Ile Gln Phe Ile Arg His Trp Arg Gln \$405\$ \$410\$ \$415\$
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 515 520 525
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 - Ser Leu Leu Leu Leu Phe Leu Phe Ile Val Ile Phe Arg Leu Leu 565 570 575
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 - Lys Ile Asp Glu Phe Glu Ser Asn Val Asn Glu Val Lys Asp Pro Tyr 725 730 735
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 - Leu Lys Met Thr Thr Tyr Gly Ala Phe Leu His Lys Gly Ser Phe Cys 850 850
- 15 Arg Asn Tyr Phe Asn Met Leu Asp Leu Leu Val Val Ala Val Ser Leu 865 870 875 880
 - Ile Ser Met Gly Leu Glu Ser Ser Ala Ile Ser Val Val Lys Ile Leu 885 890 895
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 - Lys Gly Arg Ile Lys His Leu Asp Val Val Thr Leu Leu Arg Arg 1415 1420 1425
- Ile Gln Pro Pro Leu Gly Phe Gly Lys Phe Cys Pro His Arg Val 30 1430 1435 1440

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Gln Ala Ile Ser Lys Ser Lys Leu Ser Arg Arg Trp Arg Trp Asn 50 55 60

Arg Phe Asn Arg Arg Cys Arg Ala Ala Val Lys Ser Val Thr Phe 15 65 70 75 80

Tyr Trp Leu Val Ile Val Leu Val Phe Leu Asn Thr Leu Thr Ile Ser 85 90 95

Ser Glu His Tyr Asn Gln Pro Asp Trp Leu Thr Gln Ile Gln Asp Met
100 105 110

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Glu Leu Glu Ile Met Ser Pro Leu Gly Ile Ser Val Phe Arg Cys Val 165 170 175

Arg Leu Leu Arg Ile Phe Lys Val Thr Arg His Trp Thr Ser Leu Ser 180 185 190

30 Asn Leu Val Ala Ser Leu Leu Asn Ser Met Lys Ser Ile Ala Ser Leu 195 200 205

Leu Leu Leu Phe Leu Phe Ile Ile Ile Phe Ser Leu Leu Gly Met 210 215 220

Gln Leu Phe Gly Gly Lys Phe Asn Phe Asp Glu Thr Gln Thr Lys Arg 225 230 235 240

Ser Thr Phe Asp Asn Phe Pro Gln Ala Leu Leu Thr Val Phe Gln Ile 245 250 255

5 Leu Thr Gly Glu Asp Trp Asn Ala Val Met Tyr Asp Gly Ile Met Ala
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Tyr Gly Gly Pro Ser Ser Ser Gly Met Ile Val Cys Ile Tyr Phe Ile 275 280 285

Ile Leu Phe Ile Cys Gly Asn Tyr Ile Leu Leu Asn Val Phe Leu Ala 10 290 295 300

Ile Ala Val Asp Asn Leu Ala Asp Ala Glu Ser Leu Asn Thr Ala Gln 305 310 315 320

Lys Glu Glu Ala Glu Lys Glu Arg Lys Lys Ile Ala Arg Lys Glu 325 330 335

15 Ser Leu Glu Asn Lys Lys Asn Asn Lys Pro Glu Val Asn Gln Ile Ala 340 345 350

Asn Ser Asp Asn Lys Val Thr Ile Asp Asp Tyr Arg Glu Glu Asp Glu 355 360 365

Asp Lys Asp Pro Tyr Pro Pro Cys Asp Val Pro Gly Met Val Xaa Xaa $20 \quad 370 \quad 375 \quad 380$

- Xaa Xaa Pro Cys Arg Ile Arg Val Gly Cys His Lys Leu Ile As
n His 485 490 495
- 5 His Ile Phe Thr Asn Leu Ile Leu Val Phe Ile Met Leu Ser Ser Ala 500 505 510
 - Ala Leu Ala Ala Glu Asp Pro Ile Arg Ser His Ser Phe Arg Asn Thr 515 520 525
- Ile Leu Gly Tyr Phe Asp Tyr Ala Phe Thr Ala Ile Phe Thr Val Glu 10 530 540
 - Ile Leu Leu Lys Met Thr Thr Phe Gly Ala Phe Leu His Lys Gly Ala 545 550 555 555
 - Phe Cys Arg Asn Tyr Phe Asn Leu Leu Asp Met Leu Val Val Gly Val
 565 570 575
- 15 Ser Leu Val Ser Phe Gly Ile Gln Ser Ser Ala Ile Ser Val Val Lys
 580 585 590
 - Ile Leu Arg Val Leu Arg Val Leu Arg Pro Leu Arg Ala Ile Asn Arg 595 600 605
- Ala Lys Gly Leu Lys His Val Val Gln Cys Val Phe Val Ala Ile Arg 20 610 620
 - Thr Ile Gly Asn Ile Met Ile Val Thr Thr Leu Leu Gln Phe Met Phe 625 630 635 640
 - Ala Cys Ile Gly Val Gln Leu Phe Lys Gly Lys Phe Tyr Arg Cys Thr 645 650 655
- 25 Asp Glu Ala Lys Ser Asn Pro Glu Glu Cys Arg Gly Leu Phe Ile Leu 660 665 670
 - Tyr Lys Asp Gly Asp Val Asp Ser Pro Val Val Arg Glu Arg Ile Trp 675 680 685
- Gln Asn Ser Asp Phe Asn Phe Asp Asn Val Leu Ser Ala Met Met Ala $30 \ 690 \ 695 \ 700$

Leu Phe Thr Val Ser Thr Phe Glu Gly Trp Pro Ala Leu Leu Tyr Lys
705 710 715 720

Ala Ile Asp Ser Asn Gly Glu Asn Ile Gly Pro Ile Tyr Asn His Arg
725 730 735

5 Val Glu Ile Ser Ile Phe Phe Ile Ile Tyr Ile Ile Ile Val Ala Phe 740 745 750

Phe Met Met Asn Ile Phe Val Gly Phe Val Ile Val Thr Phe Gln Glu
755 760 765

Gln Gly Glu Lys Glu Tyr Lys Asn Cys Glu Leu Asp Lys Asn Gln Arg 10 770 780

Gln Cys Val Glu Tyr Ala Leu Lys Ala Arg Pro Leu Arg Arg Tyr Ile 785 790 795 800

Pro Lys Asn Pro Tyr Gln Tyr Lys Phe Trp Tyr Val Val Asn Ser Ser 805 810 815

15 Pro Phe Glu Tyr Met Met Phe Val Leu Ile Met Leu Asn Thr Leu Cys 820 825 830

Leu Ala Met Gln His Tyr Glu Gln Ser Lys Met Phe Asn Asp Ala Met 835 840 845

Asp Ile Leu Asn Met Val Phe Thr Gly Val Phe Thr Val Glu Met Val 20 850 855 860

Leu Lys Val Ile Ala Phe Lys Pro Lys Gly Tyr Phe Ser Asp Ala Trp 865 870 870 880

Asn Thr Phe Asp Ser Leu Ile Val Ile Gly Ser Ile Ile Asp Val Ala 885 890 895

25 Leu Ser Glu Ala Lys Pro Thr Glu Ser Glu Asn Val Pro Val Pro Thr 900 905 910

Ala Thr Pro Gly Asn Ser Glu Glu Ser Asn Arg Ile Ser Ile Thr Phe 915 920 925

Phe Arg Leu Phe Arg Val Met Arg Leu Val Lys Leu Leu Ser Arg Gly 930 930 940

Glu Gly Ile Arg Thr Leu Leu Trp Thr Phe Ile Lys Ser Phe Gln Ala 945 950 955 960

Leu Pro Tyr Val Ala Leu Leu Ile Ala Met Leu Phe Phe Ile Tyr Ala 965 970 975

5 Val Ile Gly Met Gln Met Phe Gly Lys Val Ala Met Arg Asp Asn Asn 980 985 990

Gln Ile Asn Arg Asn Asn Asn Phe Gln Thr Phe Pro Gln Ala Val Leu 995 1000 1005

Leu Leu Phe Arg Cys Ala Thr Gly Glu Ala Trp Gln Glu Ile Met $10 \ 1010 \ 1015 \ 1020$

Leu Ala Cys Leu Pro Gly Lys Leu Cys Asp Pro Glu Ser Asp Tyr 1025 1030 1035

Asn Pro Gly Glu Glu Tyr Thr Cys Gly Ser Asn Phe Ala Ile Val 1040 $$ 1045 $$ 1050

15 Tyr Phe Ile Ser Phe Tyr Met Leu Cys Ala Phe Leu Ile Ile Asn 1055 1060 1065

Leu Phe Val Ala Val Ile Met Asp Asn Phe Asp Tyr Leu Thr Arg 1070 1075 1080

Asp Trp Ser Ile Leu Gly Pro His His Leu Asp Glu Phe Lys Arg $20 \ 1085 \ 1090 \ 1095$

Ile Trp Ser Glu Tyr Asp Pro Glu Ala Lys Gly Arg Ile Lys His 1100 1105 1110

Leu Asp Val Val Thr Leu Leu Arg Arg Ile Gln Pro Pro Leu Gly 1115 1120 1125

25 Phe Gly Lys Leu Cys Pro His Arg Val Ala Cys Lys Arg Leu Val 1130 1135 1140

Ala Met Asn Met Pro Leu Asn Ser Asp Gly Thr Val Met Phe Asn 1145 1150 1155

Ala Thr Leu Phe Ala Leu Val Arg Thr Ala Leu Lys Ile Lys Thr 30 1160 1165 1170

30 1385

1390

1395

Glu Gly Asn Leu Glu Gln Ala Asn Glu Glu Leu Arg Ala Val Ile 1175 1180 1185 Lys Lys Ile Trp Lys Lys Thr Ser Met Lys Leu Leu Asp Gln Val 1195 1190 5 Val Pro Pro Ala Gly Asp Asp Glu Val Thr Val Gly Lys Phe Tyr 1205 1210 1215 Ala Thr Phe Leu Ile Gln Asp Tyr Phe Arg Lys Phe Lys Lys Arg 1220 1225 1230 Lys Glu Gln Gly Leu Val Gly Lys Tyr Pro Ala Lys Asn Thr Thr 10 1235 1240 1245 Ile Ala Leu Gln Ala Gly Leu Arg Thr Leu His Asp Ile Gly Pro 1250 1255 1260 Glu Ile Arg Arg Ala Ile Ser Cys Asp Leu Gln Asp Asp Glu Pro 1265 1270 1275 15 Glu Glu Thr Lys Arg Glu Glu Glu Asp Asp Val Phe Lys Arg Asn 1280 1285 1290 Gly Ala Leu Leu Gly Asn His Val Asn His Val Asn Ser Asp Arg 1295 1300 1305 Arg Asp Ser Leu Gln Gln Thr Asn Thr His Arg Pro Leu His 20 1310 1315 1320 Val Gln Arg Pro Ser Ile Pro Pro Ala Ser Asp Thr Glu Lys Pro 1325 1330 1335 Leu Phe Pro Pro Ala Gly Asn Ser Val Cys His Asn His His Asn 1340 1345 1350 25 His Asn Ser Ile Gly Lys Gln Val Pro Thr Ser Thr Asn Ala Asn 1355 1360 Leu Asn Asn Ala Asn Met Ser Lys Ala Ala His Gly Lys Arg Pro 1370 1375 1380 Ser Ile Gly Asn Leu Glu His Val Ser Glu Asn Gly His His Ser

30 1610

Ser His Lys His Asp Arg Glu Pro Gln Arg Arg Ser Ser Val Lys 1405 1410 Arg Thr Arg Tyr Tyr Glu Thr Tyr Ile Arg Ser Asp Ser Gly Asp 1420 $5\, ext{Glu Gln}$ Leu Pro Thr Ile Cys Arg Glu Asp Pro Glu Ile His Gly 1435 Tyr Phe Arg Asp Pro His Cys Leu Gly Glu Gln Glu Tyr Phe Ser 1445 Ser Glu Glu Cys Tyr Glu Asp Asp Ser Ser Pro Thr Trp Ser Arg 10 1460 1465 Gln Asn Tyr Gly Tyr Tyr Ser Arg Tyr Pro Gly Arg Asn Ile Asp 1475 1480 Ser Glu Arg Pro Arg Gly Tyr His His Pro Gln Gly Phe Leu Glu 1490 1495 1500 15 Asp Asp Asp Ser Pro Val Cys Tyr Asp Ser Arg Arg Ser Pro Arg 1505 1510 1515 Arg Arg Leu Leu Pro Pro Thr Pro Ala Ser His Arg Arg Ser Ser 1520 1525 1530 Phe Asn Phe Glu Cys Leu Arg Arg Gln Ser Ser Gln Glu Glu Val 20 1535 1540 Pro Ser Ser Pro Ile Phe Pro His Arg Thr Ala Leu Pro Leu His 1550 1555 Leu Met Gln Gln Gln Ile Met Ala Val Ala Gly Leu Asp Ser Ser 1565 1570 25 Lys Ala Gln Lys Tyr Ser Pro Ser His Ser Thr Arg Ser Trp Ala 1580 Thr Pro Pro Ala Thr Pro Pro Tyr Arg Asp Trp Thr Pro Cys Tyr 1595 1600

Thr Pro Leu Ile Gln Val Glu Gln Ser Glu Ala Leu Asp Gln Val

1615

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	Glu	Pro 1640	Asp	Ile	Ser	Tyr	Arg 1645	Thr	Phe	Thr	Pro	Ala 1650	Ser	Leu	Thr	
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	Ala	Asp 1670	Ser	Leu	Val	Glu	Ala 1675	Val	Leu	Ile	Ser	Glu 1680	Gly	Leu	Gly	
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	Ile	Ala 1700	Asp	Ala	Cys	Asp	Leu 1705	Thr	Ile	Asp	Glu	Met 1710	Glu	Ser	Ala	
	Ala	Ser 1715	Thr	Leu	Leu	Asn	Gly 1720	Asn	Val	Arg	Pro	Arg 1725	Ala	Asn	Gly	
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	Phe	Gly 1745	Pro	Gly	Tyr	Ser	Asp 1750	Glu	Glu	Pro	Asp	Pro 1755	Gly	Arg	Asp	
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35	atg Met 1	atg a Met N	atg a Met N	Met 1	atg a Met N	atg a Met N	atg aa Met Ly	aa aa ys Ly	aa at ys Me	et Gl	ag ca ln H:	at caa is Gli	a cgg n Arg	g cag g Gli 15	g cag 1 n Gln	.66

		gcg Ala															21	L 4
5		ctt Leu															26	52
		gtc Val 50															31	LO
10		gcc Ala															35	8
15	Ser	caa Gln															40	6
		tcc Ser															45	54
20	aac Asn	ccc Pro	atc Ile 115	cga Arg	aga Arg	gcc Ala	tgc Cys	att Ile 120	agt Ser	ata Ile	gtg Val	gaa Glu	tgg Trp 125	aaa Lys	cca Pro	ttt Phe	50	2
		ata Ile 130															55	0
25	att Ile 145	tac Tyr	atc Ile	cca Pro	ttc Phe	cct Pro 150	gaa Glu	gat Asp	gat Asp	tct Ser	aat Asn 155	tca Ser	aca Thr	aat Asn	cat His	aac Asn 160	59	8
30	Leu	gaa Glu															64	.6
		ttg Leu															69	4
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45	gcc Ala	ttt Phe	cga Arg	gtg Val	ttg Leu 245	cga Arg	cca Pro	ctt Leu	cga Arg	cta Leu 250	gtg Val	tca Ser	gly aaa	gtg Val	ccc Pro 255	agt Ser	88	6

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		_	-						_				_	ttt Phe		_	1030
10														gcg Ala			1078
15	Gly													agg Arg			1126
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														cag Gln 430			1414
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	gat Asp	ccg Pro 450	gag Glu	aat Asn	gag Glu	gaa Glu	gaa Glu 455	gga Gly	gga Gly	gag Glu	gaa Glu	ggc Gly 460	aaa Lys	cga Arg	aat Asn	act Thr	1510
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		cgg Arg												1654
5		caa Gln												1702
		cga Arg 530												1750
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		ttt Phe 610												1990
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	_	cag Gln 690	-		 _			_	_	_			_	2230
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45		ctg Leu												2326

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5														gtc Val			2	2422
			-				_	_	_	_	_	_	_	aac Asn		_	2	470
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15	Glu	_		_			_					_	-	aac Asn	_		2	566
														gaa Glu 830			2	614
20														Gly 999			2	1662
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														aac Asn 910			2	854
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45	Phe	gga Gly	gct Ala	ttc Phe	ctc Leu 965	cac His	aaa Lys	gjà aaa	gcc Ala	ttc Phe 970	tgc Cys	agg Arg	aac Asn	tac Tyr	ttc Phe 975	aat Asn	3	046

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	_	_	Pro			_	atc Ile 1015		_	_				_		3187
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15	Met		Val				ctg Leu 1045									3277
			Leu				aag Lys 1060							gaa Glu		3322
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		999 Gly 1085	Asp				cct Pro 1090							tgg Trp		3412
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	_	gcc Ala 1130	Ile	-	_		gga Gly 1135							tac Tyr		3547
35			Val				atc Ile 1150							atc Ile		3592
		gct Ala 1160	Phe				aac Asn 1165									3637
40	aca Thr	ttt Phe 1175	Gln				gaa Glu 1180							gag Glu	_	3682
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		_		_				Lys				cag Gln 1215		_		3772
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			_					_	-	_	_	cag Gln 1245				3862
10												ctg Leu 1260				3907
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		gac Asp 1310										cca Pro 1320		gct Ala		4087
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												aag Lys 1365			cag Gln	4222
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		gcc Ala 1460														4	1537
10		atc Ile 1475														4	1582
15		ctg Leu 1490			-			Ile	_					tta Leu		4	1627
		ttc Phe 1505														4	1672
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		ccc Pro 1535					-	Leu	_				_		_	4	1762
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30		atg Met 1565												gct Ala		4	1852
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		gac Asp 1685						Lys								5212
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		cat His 1760					tcc Ser 1765							acc Thr		5437
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		ttc Phe 1910												tca Ser		5887
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		gac Asp 2030													tcc Ser	6247
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		aaa Lys 2105			atc Ile		gat Asp 2110	gcc Ala						gac Asp		6472
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	cata	tccat	t tt	taaa	accct	tto	cccca	aaa t	acac	ctgcg	gt co	ctggtt	cct	gttt	agctgt	7184
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Met Met Met Met Met Met Lys Lys Met Gl
n His Gl
n Arg Gl
n Gl
n 10 $$ 15

Pro Leu Ser Gly Glu Gly Pro Thr Ser Gln Pro Asn Ser Ser Lys Gln 35 40 45

Thr Val Leu Ser Trp Gln Ala Ala Ile Asp Ala Ala Arg Gln Ala Lys 50 55 60

Ala Ala Gln Thr Met Ser Thr Ser Ala Pro Pro Pro Val Gly Ser Leu 65 70 75 80

5 Ser Gln Arg Lys Arg Gln Gln Tyr Ala Lys Ser Lys Lys Gln Gly Asn 85 90 95

Ser Ser Asn Ser Arg Pro Ala Arg Ala Leu Phe Cys Leu Ser Leu Asn 100 105 110

Asn Pro Ile Arg Arg Ala Cys Ile Ser Ile Val Glu Trp Lys Pro Phe
10 115 120 125

Asp Ile Phe Ile Leu Leu Ala Ile Phe Ala Asn Cys Val Ala Leu Ala 130 135 140

Ile Tyr Ile Pro Phe Pro Glu Asp Asp Ser Asn Ser Thr Asn His Asn 145 150 155 160

15 Leu Glu Lys Val Glu Tyr Ala Phe Leu Ile Ile Phe Thr Val Glu Thr
165 170 175

Phe Leu Lys Ile Ile Ala Tyr Gly Leu Leu Leu His Pro Asn Ala Tyr
180 185 190

Val Arg Asn Gly Trp Asn Leu Leu Asp Phe Val Ile Val Ile Val Gly 20 195 200 205

Leu Phe Ser Val Ile Leu Glu Gln Leu Thr Lys Glu Thr Glu Gly Gly 210 215 220

Asn His Ser Ser Gly Lys Ser Gly Gly Phe Asp Val Lys Ala Leu Arg 225 230 235 240

25 Ala Phe Arg Val Leu Arg Pro Leu Arg Leu Val Ser Gly Val Pro Ser 245 250 255

Leu Gln Val Val Leu Asn Ser Ile Ile Lys Ala Met Val Pro Leu Leu 260 265 270

His Ile Ala Leu Leu Val Leu Phe Val Ile Ile Ile Tyr Ala Ile Ile 30 275 280 285

Gly Leu Glu Leu Phe Ile Gly Lys Met His Lys Thr Cys Phe Phe Ala 290 295 300

Asp Ser Asp Ile Val Ala Glu Glu Asp Pro Ala Pro Cys Ala Phe Ser 305 310 315 320

5 Gly Asn Gly Arg Gln Cys Thr Ala Asn Gly Thr Glu Cys Arg Ser Gly 325 330 335

Trp Val Gly Pro Asn Gly Gly Ile Thr Asn Phe Asp Asn Phe Ala Phe 340 345 350

Ala Met Leu Thr Val Phe Gln Cys Ile Thr Met Glu Gly Trp Thr Asp 355 360 365

Val Leu Tyr Trp Val Asn Asp Ala Ile Gly Trp Glu Trp Pro Trp Val 370 375 380

Tyr Phe Val Ser Leu Ile Ile Leu Gly Ser Phe Phe Val Leu Asn Leu 385 390 395 400

15 Val Leu Gly Val Leu Ser Gly Glu Phe Ser Lys Glu Arg Glu Lys Ala
405 410 415

Lys Ala Arg Gly Asp Phe Gln Lys Leu Arg Glu Lys Gln Gln Leu Glu 420 425 430

Glu Asp Leu Lys Gly Tyr Leu Asp Trp Ile Thr Gln Ala Glu Asp Ile 20 435 440 445

Asp Pro Glu Asn Glu Glu Glu Gly Glu Glu Glu Gly Lys Arg Asn Thr 450 460

Ser Met Pro Thr Ser Glu Thr Glu Ser Val Asn Thr Glu Asn Val Ser 465 470 475 480

25 Gly Glu Gly Glu Asn Arg Gly Cys Cys Gly Ser Leu Trp Cys Trp Trp 485 490 495

Arg Arg Gly Ala Ala Lys Ala Gly Pro Ser Gly Cys Arg Arg Trp $500 \hspace{1cm} 505 \hspace{1cm} 510 \hspace{1cm}$

Gly Gln Ala Ile Ser Lys Ser Lys Leu Ser Arg Arg Trp Arg Arg Trp 30 525

Asn Arg Phe Asn Arg Arg Arg Cys Arg Ala Ala Val Lys Ser Val Thr 530 535 540

Phe Tyr Trp Leu Val Ile Val Leu Val Phe Leu Asn Thr Leu Thr Ile 545 550 555 560

5 Ser Ser Glu His Tyr Asn Gln Pro Asp Trp Leu Thr Gln Ile Gln Asp
565 570 575

Ile Ala Asn Lys Val Leu Leu Ala Leu Phe Thr Cys Glu Met Leu Val 580 585 590

Lys Met Tyr Ser Leu Gly Leu Gln Ala Tyr Phe Val Ser Leu Phe Asn 600 605

Arg Phe Asp Cys Phe Val Val Cys Gly Gly Ile Thr Glu Thr Ile Leu 610 615 620

Val Glu Leu Glu Ile Met Ser Pro Leu Gly Ile Ser Val Phe Arg Cys 625 630 635 640

15 Val Arg Leu Leu Arg Ile Phe Lys Val Thr Arg His Trp Thr Ser Leu 645 650 655

Ser Asn Leu Val Ala Ser Leu Leu Asn Ser Met Lys Ser Ile Ala Ser 660 665 670

Leu Leu Leu Leu Phe Leu Phe Ile Ile Ile Phe Ser Leu Leu Gly 675 680 685

Met Gln Leu Phe Gly Gly Lys Phe Asn Phe Asp Glu Thr Gln Thr Lys 690 695 700

Arg Ser Thr Phe Asp Asn Phe Pro Gln Ala Leu Leu Thr Val Phe Gln 705 710 715 720

25 Ile Leu Thr Gly Glu Asp Trp Asn Ala Val Met Tyr Asp Gly Ile Met
725 730 735

Ala Tyr Gly Gly Pro Ser Ser Ser Gly Met Ile Val Cys Ile Tyr Phe 740 745 750

Ile Ile Leu Phe Ile Cys Gly Asn Tyr Ile Leu Leu Asn Val Phe Leu $30 \ 755 \ 760 \ 765$

- Ala Ile Ala Val Asp Asn Leu Ala Asp Ala Glu Ser Leu Asn Thr Ala 770 780
- Gln Lys Glu Glu Ala Glu Glu Lys Glu Arg Lys Lys Ile Ala Arg Lys 785 790 795 800
- 5 Glu Ser Leu Glu Asn Lys Lys Asn Asn Lys Pro Glu Val Asn Gln Ile 805 810 815
 - Ala Asn Ser Asp Asn Lys Val Thr Ile Asp Asp Tyr Arg Glu Glu Asp 820 825 830
- Glu Asp Lys Asp Pro Tyr Pro Pro Cys Asp Val Pro Val Gly Glu Glu 10 835 840 845
 - Glu Glu Glu Glu Glu Asp Glu Pro Glu Val Pro Ala Gly Pro Arg 850 855 860
 - Pro Arg Arg Ile Ser Glu Leu Asn Met Lys Glu Lys Ile Ala Pro Ile 865 870 875 888
- 15 Pro Glu Gly Ser Ala Phe Phe Ile Leu Ser Lys Thr Asn Pro Ile Arg 885 890 895
 - Val Gly Cys His Lys Leu Ile Asn His His Ile Phe Thr Asn Leu Ile 900 905 910
- Leu Val Phe Ile Met Leu Ser Ser Ala Ala Leu Ala Ala Glu Asp Pro 915 920 925
 - Ile Arg Ser His Ser Phe Arg Asn Thr Ile Leu Gly Tyr Phe Asp Tyr 930 935 940
 - Ala Phe Thr Ala Ile Phe Thr Val Glu Ile Leu Leu Lys Met Thr Thr 945 950 955 960
- 25 Phe Gly Ala Phe Leu His Lys Gly Ala Phe Cys Arg Asn Tyr Phe Asn 965 970 975
 - Leu Leu Asp Met Leu Val Val Gly Val Ser Leu Val Ser Phe Gly Ile 980 985 990
- Gln Ser Ser Ala Ile Ser Val Val Lys Ile Leu Arg Val Leu Arg Val 30 995 1000 1005

1225

1230

Leu Arg Pro Leu Arg Ala Ile Asn Arg Ala Lys Gly Leu Lys His 1015 1020 1010 Val Val Gln Cys Val Phe Val Ala Ile Arg Thr Ile Gly Asn Ile 1030 1025 1035 5 Met Ile $\,$ Val Thr Thr Leu Leu $\,$ Gln $\,$ Phe $\,$ Met $\,$ Phe $\,$ Ala $\,$ Cys $\,$ Ile $\,$ Gly 1045 1040 1050 Val Gln Leu Phe Lys Gly Lys Phe Tyr Arg Cys Thr Asp Glu Ala 1055 1060 1065 Lys Ser Asn Pro Glu Glu Cys Arg Gly Leu Phe Ile Leu Tyr Lys 10 1070 1075 1080 Asp Gly Asp Val Asp Ser Pro Val Val Arg Glu Arg Ile Trp Gln 1085 1090 1095 Asn Ser Asp Phe Asn Phe Asp Asn Val Leu Ser Ala Met Met Ala 1100 1105 1110 15 Leu Phe Thr Val Ser Thr Phe Glu Gly Trp Pro Ala Leu Leu Tyr 1115 1120 1125 Lys Ala Ile Asp Ser Asn Gly Glu Asn Ile Gly Pro Ile Tyr Asn 1130 1135 1140 His Arg Val Glu Ile Ser Ile Phe Phe Ile Ile Tyr Ile Ile Ile 20 1145 1150 1155 Val Ala Phe Phe Met Met Asn Ile Phe Val Gly Phe Val Ile Val 1160 1165 1170 Thr Phe Gln Glu Gln Gly Glu Lys Glu Tyr Lys Asn Cys Glu Leu 1175 1180 1185 25 Asp Lys Asn Gln Arg Gln Cys Val Glu Tyr Ala Leu Lys Ala Arg 1190 1195 1200 Pro Leu Arg Arg Tyr Ile Pro Lys Asn Pro Tyr Gln Tyr Lys Phe 1205 1210 1215 Trp Tyr Val Val Asn Ser Ser Pro Phe Glu Tyr Met Met Phe Val

1450

1455

Leu Ile Met Leu Asn Thr Leu Cys Leu Ala Met Gln His Tyr Glu 1235 1240 1245 Gln Ser Lys Met Phe Asn Asp Ala Met Asp Ile Leu Asn Met Val 1255 1250 1260 5 Phe Thr Gly Val Phe Thr Val Glu Met Val Leu Lys Val Ile Ala 1265 1270 1275 Phe Lys Pro Lys Gly Tyr Phe Ser Asp Ala Trp Asn Thr Phe Asp 1280 1285 1290 Ser Leu Ile Val Ile Gly Ser Ile Ile Asp Val Ala Leu Ser Glu 10 1295 1300 1305 Ala Asp Pro Thr Glu Ser Glu Asn Val Pro Val Pro Thr Ala Thr 1310 1315 1320 Pro Gly Asn Ser Glu Glu Ser Asn Arg Ile Ser Ile Thr Phe Phe 1325 1330 1335 $15\,\mathrm{Arg}$ Leu Phe Arg Val Met Arg Leu Val Lys Leu Leu Ser Arg Gly 1340 1345 1350 Glu Gly Ile Arg Thr Leu Leu Trp Thr Phe Ile Lys Ser Phe Gln 1355 1360 1365 Ala Leu Pro Tyr Val Ala Leu Leu Ile Ala Met Leu Phe Phe Ile 20 1370 1375 1380 Tyr Ala Val Ile Gly Met Gln Met Phe Gly Lys Val Ala Met Arg 1385 1390 1395 Asp Asn Asn Gln Ile Asn Arg Asn Asn Phe Gln Thr Phe Pro 1400 1405 1410 25 Gln Ala Val Leu Leu Phe Arg Cys Ala Thr Gly Glu Ala Trp 1415 1420 1425 Gln Glu Ile Met Leu Ala Cys Leu Pro Gly Lys Leu Cys Asp Pro 1430 1435 1440 Glu Ser Asp Tyr Asn Pro Gly Glu Glu Tyr Thr Cys Gly Ser Asn Phe Ala Ile Val Tyr Phe Ile Ser Phe Tyr Met Leu Cys Ala Phe 1460 1465 1470

Leu Ile Ile Asn Leu Phe Val Ala Val Ile Met Asp Asn Phe Asp 1475 1480 1485

5 Tyr Leu Thr Arg Asp Trp Ser Ile Leu Gly Pro His His Leu Asp 1490 1495 1500

Glu Phe Lys Arg Ile Trp Ser Glu Tyr Asp Pro Glu Ala Lys Gly 1505 1510 1515

Arg Ile Lys His Leu Asp Val Val Thr Leu Leu Arg Arg Ile Gln 1520 1525 1530

Pro Pro Leu Gly Phe Gly Lys Leu Cys Pro His Arg Val Ala Cys 1535 1540 1545

Lys Arg Leu Val Ala Met Asn Met Pro Leu Asn Ser Asp Gly Thr 1550 1560

15 Val Met Phe Asn Ala Thr Leu Phe Ala Leu Val Arg Thr Ala Leu 1565 1570 1575

Lys Ile Lys Thr Glu Gly Asn Leu Glu Gln Ala Asn Glu Glu Leu 1580 . 1585 1590

Arg Ala Val Ile Lys Lys Ile Trp Lys Lys Thr Ser Met Lys Leu $20 \ 1595 \ 1600 \ 1605$

Leu Asp Gln Val Val Pro Pro Ala Gly Asp Asp Glu Val Thr Val
. 1610 1615 1620

Glý Lys Phe Tyr Ala Thr Phe Leu Ile Gln Asp Tyr Phe Arg Lys 1625 1630 1635

25 Phe Lys Lys Arg Lys Glu Gln Gly Leu Val Gly Lys Tyr Pro Ala 1640 1645 1650

Lys Asn Thr Thr Ile Ala Leu Gln Ala Gly Leu Arg Thr Leu His 1655 1660 1665

Asp Ile Gly Pro Glu Ile Arg Arg Ala Ile Ser Cys Asp Leu Gln 1670 1675 1680

Asp Asp Glu Pro Glu Glu Thr Lys Arg Glu Glu Glu Asp Asp Val 1685 1690 Phe Lys Arg Asn Gly Ala Leu Leu Gly Asn His Val Asn His Val 1705 1700 ${f 5}$ Asn Ser Asp Arg Arg Asp Ser Leu Gln Gln Thr Asn Thr Thr His 1720 Arg Pro Leu His Val Gln Arg Pro Ser Ile Pro Pro Ala Ser Asp 1730 Thr Glu Lys Pro Leu Phe Pro Pro Ala Gly Asn Ser Val Cys His 10 1745 1750 1755 Asn His Asn His Asn Ser Ile Gly Lys Gln Val Pro Thr Ser 1765 1770 Thr Asn Ala Asn Leu Asn Asn Ala Asn Met Ser Lys Ala Ala His 1775 1780 1785 $15\,\mathrm{Gly}$ Lys Arg Pro Ser Ile Gly Asn Leu Glu His Val Ser Glu Asn 1790 1795 1800 Gly His His Ser Ser His Lys His Asp Arg Glu Pro Gln Arg Arg 1805 1810 1815 Ser Ser Val Lys Arg Thr Arg Tyr Tyr Glu Thr Tyr Ile Arg Ser 20 1820 1825 1830 Asp Ser Gly Asp Glu Gln Leu Pro Thr Ile Cys Arg Glu Asp Pro 1835 1840 1845 Glu Ile His Gly Tyr Phe Arg Asp Pro His Cys Leu Gly Glu Gln 1850 1855 1860 $25\,\mathrm{Glu}$ Tyr Phe Ser Ser Glu Glu Cys Tyr Glu Asp Asp Ser Ser Pro 1865 1870 1875 Thr Trp Ser Arg Gln Asn Tyr Gly Tyr Tyr Ser Arg Tyr Pro Gly 1880 1885 1890 Arg Asn Ile Asp Ser Glu Arg Pro Arg Gly Tyr His His Pro Gln

30 1895 1900 1905

-221-Gly Phe Leu Glu Asp Asp Ser Pro Val Cys Tyr Asp Ser Arg 1915 Arg Ser Pro Arg Arg Leu Leu Pro Pro Thr Pro Ala Ser His 1930 1935 1925 ${f 5}$ Arg Arg Ser Ser Phe Asn Phe Glu Cys Leu Arg Arg Gln Ser Ser 1945 1940 1950 Gln Glu Glu Val Pro Ser Ser Pro Ile Phe Pro His Arg Thr Ala 1955 1960 Leu Pro Leu His Leu Met Gln Gln Gln Ile Met Ala Val Ala Gly 10 1970 1975 1980 Leu Asp Ser Ser Lys Ala Gln Lys Tyr Ser Pro Ser His Ser Thr 1985 1990 1995 Arg Ser Trp Ala Thr Pro Pro Ala Thr Pro Pro Tyr Arg Asp Trp 2000 2005 2010

15 Thr Pro Cys Tyr Thr Pro Leu Ile Gln Val Glu Gln Ser Glu Ala 2015 2020 2025

Leu Asp Gln Val Asn Gly Ser Leu Pro Ser Leu His Arg Ser Ser 2030 2035 2040

Trp Tyr Thr Asp Glu Pro Asp Ile Ser Tyr Arg Thr Phe Thr Pro 20 2045 2050 2055

Ala Ser Leu Thr Val Pro Ser Ser Phe Arg Asn Lys Asn Ser Asp 2060 2065 2070

Lys Gln Arg Ser Ala Asp Ser Leu Val Glu Ala Val Leu Ile Ser 2075 2080 2085

 $25\,\text{Glu}$ Gly Leu Gly Arg Tyr Ala Arg Asp Pro Lys Phe Val Ser Ala 2090 2095 2100

Thr Lys His Glu Ile Ala Asp Ala Cys Asp Leu Thr Ile Asp Glu 2105 2110 2115

Met Glu Ser Ala Ala Ser Thr Leu Leu Asn Gly Asn Val Arg Pro $30\,$ 2120 2125 2130

Arg Ala Asn Gly Asp Val Gly Pro Leu Ser His Arg Gln Asp Tyr 2135 2140 2145 Glu Leu Gln Asp Phe Gly Pro Gly Tyr Ser Asp Glu Glu Pro Asp 2150 2155 2160 5 Pro Gly Arg Asp Glu Glu Asp Leu Ala Asp Glu Met Ile Cys Ile 2165 2170 Thr Thr Leu 2180 <210> 19 10 <211> 7193 <212> DNA <213> Homo sapiens <220> <221> gene 15 <222> (1)..(7193) <220> <221> CDS <222> (119)..(6664) <400> 19 20 agaataaggg cagggaccgc ggctcctatc tcttggtgat ccccttcccc attccgcccc 60 egecteaaeg eccageaeag tgeeetgeae acagtagteg etcaataaat gttegtgg 118 atg atg atg atg atg atg aaa aaa atg cag cat caa cgg cag cag 166 Met Met Met Met Met Met Lys Lys Met Gln His Gln Arg Gln Gln 1 5 $25\,$ caa gcg gac cac gcg aac gag gca aac tat gca aga ggc acc aga ctt 214 Gln Ala Asp His Ala Asn Glu Ala Asn Tyr Ala Arg Gly Thr Arg Leu 20 cct ctt tct ggt gaa gga cca act tct cag ccg aat agc tcc aag caa 262 Pro Leu Ser Gly Glu Gly Pro Thr Ser Gln Pro Asn Ser Ser Lys Gln 30 35 act gtc ctg tct tgg caa gct gca atc gat gct gct aga cag gcc aag 310 Thr Val Leu Ser Trp Gln Ala Ala Ile Asp Ala Ala Arg Gln Ala Lys 50 55 get gee caa act atg age ace tet gea eee eea eet gta gga tet ete 358 35 Ala Ala Gln Thr Met Ser Thr Ser Ala Pro Pro Pro Val Gly Ser Leu 65 70 tcc caa aga aaa cgt cag caa tac gcc aag agc aaa aaa cag ggt aac 406 Ser Gln Arg Lys Arg Gln Gln Tyr Ala Lys Ser Lys Lys Gln Gly Asn 85 90 $40\; ext{tcg}$ tcc aac agc cga cct gcc cgc gcc ctt ttc tgt tta tca ctc aat 454 Ser Ser Asn Ser Arg Pro Ala Arg Ala Leu Phe Cys Leu Ser Leu Asn 100 105

		ccc Pro		-	_	-	_		_			_					502
5		ata Ile 130															550
		tac Tyr					_	_	_								598
10		gaa Glu															646
15	Phe	ttg Leu															694
	_	agg Arg						_	_		_		_		_		742
20		ttt Phe 210															790
		cac His															838
25		ttt Phe															886
30		caa Gln									_	_	_				934
		ata Ile															982
35	gga Gly	ttg Leu 290	gaa Glu	ctt Leu	ttt Phe	att Ile	gga Gly 295	aaa Lys	atg Met	cac His	aaa Lys	aca Thr 300	tgt Cys	ttt Phe	ttt Phe	gct Ala	1030
		tca Ser															1078
40		aat Asn															1126
45		gtt Val	-	_								_			_		1174

		atg Met															1222
5		ctc Leu 370															1270
		ttt Phe											_			_	1318
10		ctt Leu															1366
15	Lys	gca Ala			_		_	_				_	_	_	_		1414
		gat Asp		_			_	_					_		_		1462
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25		gaa Glu															1606
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40	tcc Ser	tct Ser	gag Glu	cac His	tac Tyr 565	aat Asn	cag Gln	cca Pro	gat Asp	tgg Trp 570	ttg Leu	aca Thr	cag Gln	att Ile	caa Gln 575	gat Asp	1846
45	Ile	gcc Ala	aac Asn	aaa Lys 580	gtc Val	ctc Leu	ttg Leu	gct Ala	ctg Leu 585	ttc Phe	acc Thr	tgc Cys	gag Glu	atg Met 590	ctg Leu	gta Val	1894

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5				tgc Cys													1990
				gaa Glu													2038
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	_	_		ctg Leu										_			2182
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				ttt Phe													2278
25				ggc													2326
30				ggc Gly 740									-				2374
				ttc Phe			-					_		_		_	2422
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				gaa Glu													2518
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45	gcc Ala	aac Asn	agt Ser	gac Asp 820	aac Asn	aag Lys	gtt Val	aca Thr	att Ile 825	gat Asp	gac Asp	tat Tyr	aga Arg	gaa Glu 830	gag Glu	gat Asp	2614

				gac Asp									2662
5				gag Glu									2710
				atc Ile									2758
10				agc Ser									2806
15	_		_	cac His 900	_								2854
				atc Ile									2902
20				cac His									2950
				gcc Ala									2998
25				ttc Phe									3046
30				atg Met 980									3094
								Lys			l L	gg gtc rg Val	3142
35			Pro	c cto Dei			e As			rs G			3187
		gtc Val 1025	Glr	g tgo 1 Cys			L Al			nr I			3232
40			Va]	c act L Thr			ı G			ne Al			3277
45	Val		Let	g tto 1 Phe			s Ph			s Tl			3322

	aa Ly	a agt s Ser 107	. As	ac co sn Pr	t ga o Gl	a ga u Gl	a tgc u Cys 107	Ar	g gg g Gl	a ct y Le	t tt u Ph	c ato e Ile 108	Le	c ta u Ty	c aa r Ly	g s	3367
	ga 5 As	t ggg p Gly 108	AS	ıt gt sp Va	t ga l As	c agt p Sei	t cct r Pro 1090	Va	g gt	c cg l Ar	t ga g Gl	a cgg u Arg 109	I1	c tg e Tr	g caa p Gli	a n	3412
	AS.	110	0	p Pn	e As:	n Phe	1105	Ası 5	n Val	l Le	u Se:	t gct r Ala 111	Me	t Me	t Ala	₹.	3457
1	O cto Lei	c ttc u Phe 111	T 11	a gt r Va	c tco l Se:	c acg	ttt Phe 1120	GI	r GJ ⁷ 3 330	c tgg / Trj	g cci p Pro	t gcg D Ala 112	Le	g ct	g tat u Tyr	3	3502
1	шу.	a gcc s Ala 113	T T.	c gad e Asj	c tog p Sei	g aat Asn	gga Gly 1135	GI	g aac ı Asr	ato Ile	ggo Gly	C CCa / Pro 1140	11ϵ	c tag	c aac r Asn	; 1	3547
	Cac	c cgc s Arg 114!	va.	g gag l Gli	g ato 1 Ile	tcc Ser	atc Ile 1150	Phe	ttc Phe	ato : Ile	ato Ele	tac Tyr 1155	$I1\epsilon$		c att		3592
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		Phe	GII	g gaa 1 Glu	caa Gln	gga Gly	gaa Glu 1180	Lys	gag Glu	tat Tyr	aag Lys	aac Asn 1185	Cys	gag Glu	ı ctg ı Leu		3682
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5		gac Asp 1310												gct Ala		4087
		999 Gly 1325										atc Ile 1335		ttt Phe		4132
10		ctt Leu 1340												agg Arg		4177
15	Glu	ggc Gly 1355												ttt Phe		4222
		ctc Leu 1370										ctg Leu 1380		ttc Phe		4267
20		gcg Ala 1385												atg Met	aga Arg	4312
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30	Gln	gag Glu 1430										ctc Leu 1440		gac Asp		4447
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	_	atc Ile 1475			_			_	_		-	_			_	4582
40	tat Tyr	ctg Leu 1490	acc Thr	cgg Arg	gac Asp	tgg Trp	tct Ser 1495	att Ile	ttg Leu	Gly aaa	cct Pro	cac His 1500	cat His	tta Leu	gat Asp	4627
45	Glu	ttc Phe 1505														4672

		ata Ile 1520														4717
5		ccc Pro 1535										agg Arg 1545				4762
	_	aga Arg 1550		-	_	_						_	_	gly aaa		4807
10		atg Met 1565										cga Arg 1575		gct Ala		4852
15	Lys	atc Ile 1580												gaa Glu		4897
		gct Ala 1595							_			_	-	aaa Lys		4942
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30	Lys	aac Asn 1655					cta Leu 1660					agg Arg 1665		ctg Leu		5122
	_	att Ile 1670										tgt Cys 1680			caa Gln	5167
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		aaa Lys 1700										gtc Val 1710				5257
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45	cgt Arg	ccc Pro 1730	ctg Leu	cat His	gtc Val	caa Gln	agg Arg 1735	cct Pro	tca Ser	att Ile	cca Pro	cct Pro 1740	gca Ala	agt Ser	gat Asp	5347

		gag Glu 1745						Pro								5392
5	aac Asn	cat His 1760	cat His	aac Asn	cat His	aat Asn	tcc Ser 1765	ata Ile	gga Gly	aag Lys	caa Gln	gtt Val 1770	ccc Pro	acc Thr	tca Ser	5437
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10	gga Gly	aag Lys 1790	cgg Arg	ccc Pro	agc Ser	att Ile	999 Gly 1795	aac Asn	ctt Leu	gag Glu	cat His	gtg Val 1800			aat Asn	5527
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20		tca Ser 1835					ctc Leu 1840									5662
		ata Ile 1850					agg Arg 1855							gag Glu		5707
25		tat Tyr 1865					gaa Glu 1870							tcg Ser		5752
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		tcg Ser 2000		_				_						gac Asp	-	6157
10		ccg Pro 2015														6202
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		tac Tyr 2045												aca Thr		6292
20		agc Ser 2060												agc Ser	_	6337
		cag Gln 2075												ata Ile		6382
25	gaa Glu	ggc Gly 2090	ttg Leu	gga Gly	cgc Arg	tat Tyr	gca Ala 2095	agg Arg	gac Asp	cca Pro	aaa Lys	ttt Phe 2100	gtg Val	tca Ser	gca Ala	6427
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		gag Glu 2120														6517
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		cta Leu 2150														6607
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Pro Leu Ser Gly Glu Gly Pro Thr Ser Gln Pro Asn Ser Ser Lys Gln 20 35 40 45

Thr Val Leu Ser Trp Gln Ala Ala Ile Asp Ala Ala Arg Gln Ala Lys 50 55 60

Ala Ala Gln Thr Met Ser Thr Ser Ala Pro Pro Pro Val Gly Ser Leu 65 70 75 80

25 Ser Gln Arg Lys Arg Gln Gln Tyr Ala Lys Ser Lys Lys Gln Gly Asn
85 90 95

Ser Ser Asn Ser Arg Pro Ala Arg Ala Leu Phe Cys Leu Ser Leu Asn 100 105 110

Asn Pro Ile Arg Arg Ala Cys Ile Ser Ile Val Glu Trp Lys Pro Phe 30 115 120 125

Asp Ile Phe Ile Leu Leu Ala Ile Phe Ala Asn Cys Val Ala Leu Ala 130 135 140

Ile Tyr Ile Pro Phe Pro Glu Asp Asp Ser Asn Ser Thr Asn His Asn 145 150 155 160

Leu Glu Lys Val Glu Tyr Ala Phe Leu Ile Ile Phe Thr Val Glu Thr
165 170 175

5 Phe Leu Lys Ile Ile Ala Tyr Gly Leu Leu His Pro Asn Ala Tyr
180 185 190

Val Arg Asn Gly Trp Asn Leu Leu Asp Phe Val Ile Val Ile Val Gly
195 200 205

Leu Phe Ser Val Ile Leu Glu Gln Leu Thr Lys Glu Thr Glu Gly Gly $10 \hspace{1cm} 210 \hspace{1cm} 215 \hspace{1cm} 220 \hspace{1cm}$

Asn His Ser Ser Gly Lys Ser Gly Gly Phe Asp Val Lys Ala Leu Arg 225 230 235

Ala Phe Arg Val Leu Arg Pro Leu Arg Leu Val Ser Gly Val Pro Ser 245 250 255

15 Leu Gln Val Val Leu Asn Ser Ile Ile Lys Ala Met Val Pro Leu Leu 260 265 270

His Ile Ala Leu Leu Val Leu Phe Val Ile Ile Ile Tyr Ala Ile Ile 275 280 285

Gly Leu Glu Leu Phe Ile Gly Lys Met His Lys Thr Cys Phe Phe Ala $20 \hspace{1.5cm} 290 \hspace{1.5cm} 295 \hspace{1.5cm} 300 \hspace{1.5cm}$

Asp Ser Asp Ile Val Ala Glu Glu Asp Pro Ala Pro Cys Ala Phe Ser 305 310 315 320

Gly Asn Gly Arg Gln Cys Thr Ala Asn Gly Thr Glu Cys Arg Ser Gly \$325\$ \$330 \$35

25 Trp Val Gly Pro Asn Gly Gly Ile Thr Asn Phe Asp Asn Phe Ala Phe 340 345 350

Ala Met Leu Thr Val Phe Gln Cys Ile Thr Met Glu Gly Trp Thr Asp 355 360 365

Val Leu Tyr Trp Val Asn Asp Ala Ile Gly Trp Glu Trp Pro Trp Val 30 370 380

Tyr Phe Val Ser Leu Ile Ile Leu Gly Ser Phe Phe Val Leu Asn Leu 385 390 395 400

Val Leu Gly Val Leu Ser Gly Glu Phe Ser Lys Glu Arg Glu Lys Ala 405 410 415

5 Lys Ala Arg Gly Asp Phe Gln Lys Leu Arg Glu Lys Gln Gln Leu Glu
420 425 430

Glu Asp Leu Lys Gly Tyr Leu Asp Trp Ile Thr Gln Ala Glu Asp Ile 435 440 445

Asp Pro Glu Asn Glu Glu Glu Gly Glu Glu Glu Gly Lys Arg Asn Thr $10 \hspace{0.4cm} 450 \hspace{0.4cm} 460$

Ser Met Pro Thr Ser Glu Thr Glu Ser Val Asn Thr Glu Asn Val Ser 465 470 475 480

Gly Glu Gly Glu Asn Arg Gly Cys Cys Gly Ser Leu Trp Cys Trp Trp 485 490 495

15 Arg Arg Gly Ala Ala Lys Ala Gly Pro Ser Gly Cys Arg Arg Trp 500 505 510

Gly Gln Ala Ile Ser Lys Ser Lys Leu Ser Arg Arg Trp Arg Trp 515 520 525

Asn Arg Phe Asn Arg Arg Cys Arg Ala Ala Val Lys Ser Val Thr 20 530 540

Phe Tyr Trp Leu Val Ile Val Leu Val Phe Leu Asn Thr Leu Thr Ile 545 550 560

Ser Ser Glu His Tyr Asn Gln Pro Asp Trp Leu Thr Gln Ile Gln Asp 565 570 575

25 Ile Ala Asn Lys Val Leu Leu Ala Leu Phe Thr Cys Glu Met Leu Val 580 585 590

Lys Met Tyr Ser Leu Gly Leu Gln Ala Tyr Phe Val Ser Leu Phe Asn 595 600 605

Arg Phe Asp Cys Phe Val Val Cys Gly Gly Ile Thr Glu Thr Ile Leu $30 \ \ \, 610 \ \ \, \, 615 \ \ \, \, 620$

Val Glu Leu Glu Ile Met Ser Pro Leu Gly Ile Ser Val Phe Arg Cys 625 630 635

Val Arg Leu Leu Arg Ile Phe Lys Val Thr Arg His Trp Thr Ser Leu 645 650 655

5 Ser Asn Leu Val Ala Ser Leu Leu Asn Ser Met Lys Ser Ile Ala Ser 660 670

Leu Leu Leu Leu Phe Leu Phe Ile Ile Ile Phe Ser Leu Leu Gly 675 680 685

Met Gln Leu Phe Gly Gly Lys Phe Asn Phe Asp Glu Thr Gln Thr Lys 10 690 695 700

Arg Ser Thr Phe Asp Asn Phe Pro Gln Ala Leu Leu Thr Val Phe Gln 705 710 715 720

Ile Leu Thr Gly Glu Asp Trp Asn Ala Val Met Tyr Asp Gly Ile Met 725 730 735

15 Ala Tyr Gly Gly Pro Ser Ser Ser Gly Met Ile Val Cys Ile Tyr Phe
740 745 750

Ile Ile Leu Phe Ile Cys Gly Asn Tyr Ile Leu Leu Asn Val Phe Leu 755 760 765

Ala Ile Ala Val Asp Asn Leu Ala Asp Ala Glu Ser Leu Asn Thr Ala 20 770 780

Gln Lys Glu Glu Ala Glu Glu Lys Glu Arg Lys Lys Ile Ala Arg Lys 785 790 795 800

Glu Ser Leu Glu Asn Lys Lys Asn Asn Lys Pro Glu Val Asn Gln Ile 805 810 815

25 Ala Asn Ser Asp Asn Lys Val Thr Ile Asp Asp Tyr Arg Glu Glu Asp 820 825 830

Glu Asp Lys Asp Pro Tyr Pro Pro Cys Asp Val Pro Val Gly Glu Glu 835 840 845

Glu Glu Glu Glu Glu Glu Asp Glu Pro Glu Val Pro Ala Gly Pro Arg 30 850 855 860

Pro Arg Arg Ile Ser Glu Leu Asn Met Lys Glu Lys Ile Ala Pro Ile 865 870 885 886

Pro Glu Gly Ser Ala Phe Phe Ile Leu Ser Lys Thr Asn Pro Ile Arg 885 890 895

5 Val Gly Cys His Lys Leu Ile Asn His His Ile Phe Thr Asn Leu Ile
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Leu Val Phe Ile Met Leu Ser Ser Ala Ala Leu Ala Ala Glu Asp Pro 915 920 925

Ile Arg Ser His Ser Phe Arg Asn Thr Ile Leu Gly Tyr Phe Asp Tyr 10 930 935 940

Ala Phe Thr Ala Ile Phe Thr Val Glu Ile Leu Leu Lys Met Thr Thr 945 950 955 960

Phe Gly Ala Phe Leu His Lys Gly Ala Phe Cys Arg Asn Tyr Phe Asn 965 970 975

15 Leu Leu Asp Met Leu Val Val Gly Val Ser Leu Val Ser Phe Gly Ile 980 985 990

Gln Ser Ser Ala Ile Ser Val Val Lys Ile Leu Arg Val Leu Arg Val 995 1000 1005

Leu Arg Pro Leu Arg Ala Ile Asn Arg Ala Lys Gly Leu Lys His 20 1010 1015 1020

Val Val Gln Cys Val Phe Val Ala Ile Arg Thr Ile Gly Asn Ile 1025 1030 1035

Met Ile Val Thr Thr Leu Leu Gln Phe Met Phe Ala Cys Ile Gly 1040 1045 1050

25 Val Gln Leu Phe Lys Gly Lys Phe Tyr Arg Cys Thr Asp Glu Ala 1055 1060 1065

Lys Ser Asn Pro Glu Glu Cys Arg Gly Leu Phe Ile Leu Tyr Lys 1070 1075 1080

Asp Gly Asp Val Asp Ser Pro Val Val Arg Glu Arg Ile Trp Gln 30 1085 1090 1095

Asn Ser Asp Phe Asn Phe Asp Asn Val Leu Ser Ala Met Met Ala 1100 1105 Leu Phe Thr Val Ser Thr Phe Glu Gly Trp Pro Ala Leu Leu Tyr 1115 $5\,\mathrm{Lys}$ Ala Ile Asp Ser Asn Gly Glu Asn Ile Gly Pro Ile Tyr Asn 1130 1135 His Arg Val Glu Ile Ser Ile Phe Phe Ile Ile Tyr Ile Ile Ile , 1150 1145 1155 Val Ala Phe Phe Met Met Asn Ile Phe Val Gly Phe Val Ile Val 10 1160 1165 1170 Thr Phe Gln Glu Gln Gly Glu Lys Glu Tyr Lys Asn Cys Glu Leu 1175 1180 Asp Lys Asn Gln Arg Gln Cys Val Glu Tyr Ala Leu Lys Ala Arg 1190 1195 1200 15 Pro Leu Arg Arg Tyr Ile Pro Lys Asn Pro Tyr Gln Tyr Lys Phe 1205 1210 1215 Trp Tyr Val Val Asn Ser Ser Pro Phe Glu Tyr Met Met Phe Val 1220 1225 1230 Leu Ile Met Leu Asn Thr Leu Cys Leu Ala Met Gln His Tyr Glu 20 1235 1240 1245 Gln Ser Lys Met Phe Asn Asp Ala Met Asp Ile Leu Asn Met Val 1250 1255 1260 Phe Thr Gly Val Phe Thr Val Glu Met Val Leu Lys Val Ile Ala 1265 1270 1275 25 Phe Lys Pro Lys Gly Tyr Phe Ser Asp Ala Trp Asn Thr Phe Asp 1280 1285 1290 Ser Leu Ile Val Ile Gly Ser Ile Ile Asp Val Ala Leu Ser Glu 1295 1300 1305

Ala Asp Pro Thr Glu Ser Glu Asn Val Pro Val Pro Thr Ala Thr

1315 1320

Pro Gly Asn Ser Glu Glu Ser Asn Arg Ile Ser Ile Thr Phe Phe 1325 1330 1335

Arg Leu Phe Arg Val Met Arg Leu Val Lys Leu Leu Ser Arg Gly 1340 1345 1350

5 Glu Gly Ile Arg Thr Leu Leu Trp Thr Phe Ile Lys Ser Phe Gln 1355 1360 1365

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Tyr Ala Val Ile Gly Met Gln Met Phe Gly Lys Val Ala Met Arg 1385 1390 1395

Asp Asn Asn Gln Ile Asn Arg Asn Asn Asn Phe Gln Thr Phe Pro 1400 1405 1410

Gln Ala Val Leu Leu Leu Phe Arg Cys Ala Thr Gly Glu Ala Trp 1415 1420 1425

15 Gln Glu Ile Met Leu Ala Cys Leu Pro Gly Lys Leu Cys Asp Pro 1430 1435 1440

Glu Ser Asp Tyr Asn Pro Gly Glu Glu Tyr Thr Cys Gly Ser Asn 1445 1450 1455

Phe Ala Ile Val Tyr Phe Ile Ser Phe Tyr Met Leu Cys Ala Phe 20 1460 1465 1470

Leu Ile Ile Asn Leu Phe Val Ala Val Ile Met Asp Asn Phe Asp 1475 1480 1485

Tyr Leu Thr Arg Asp Trp Ser Ile Leu Gly Pro His His Leu Asp 1490 1495 1500

25 Glu Phe Lys Arg Ile Trp Ser Glu Tyr Asp Pro Glu Ala Lys Gly
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Arg Ile Lys His Leu Asp Val Val Thr Leu Leu Arg Arg Ile Gln
1520 1525 1530

Pro Pro Leu Gly Phe Gly Lys Leu Cys Pro His Arg Val Ala Cys 30 1535 1540 1545

Lys Arg Leu Val Ala Met Asn Met Pro Leu Asn Ser Asp Gly Thr 1550 1555 Val Met Phe Asn Ala Thr Leu Phe Ala Leu Val Arg Thr Ala Leu 1565 1570 5 Lys Ile Lys Thr Glu Gly Asn Leu Glu Gln Ala Asn Glu Glu Leu 1580 1585 Arg Ala Val Ile Lys Lys Ile Trp Lys Lys Thr Ser Met Lys Leu 1595 1600 Leu Asp Gln Val Val Pro Pro Ala Gly Asp Asp Glu Val Thr Val 10 1610 1615 1620 Gly Lys Phe Tyr Ala Thr Phe Leu Ile Gln Asp Tyr Phe Arg Lys 1625 1630 1635 Phe Lys Lys Arg Lys Glu Gln Gly Leu Val Gly Lys Tyr Pro Ala 1645 1640 1650 $15\,\mathrm{Lys}$ Asn Thr Thr Ile Ala Leu Gln Ala Gly Leu Arg Thr Leu His 1655 1660 Asp Ile Gly Pro Glu Ile Arg Arg Ala Ile Ser Cys Asp Leu Gln 1670 1675 Asp Asp Glu Pro Glu Glu Thr Lys Arg Glu Glu Glu Asp Asp Val 20 1685 1690 Phe Lys Arg Asn Gly Ala Leu Leu Gly Asn His Val Asn His Val 1700 1705 Asn Ser Asp Arg Arg Asp Ser Leu Gln Gln Thr Asn Thr Thr His 1715 1720 25 Arg Pro Leu His Val Gln Arg Pro Ser Ile Pro Pro Ala Ser Asp 1730 1735 Thr Glu Lys Pro Leu Phe Pro Pro Ala Gly Asn Ser Val Cys His 1745 1750 1755

Asn His His Asn His Asn Ser Ile Gly Lys Gln Val Pro Thr Ser

1765

Thr Asn Ala Asn Leu Asn Asn Ala Asn Met Ser Lys Ala Ala His 1775 1780 1785

Gly Lys Arg Pro Ser Ile Gly Asn Leu Glu His Val Ser Glu Asn 1790 1800

5 Gly His His Ser Ser His Lys His Asp Arg Glu Pro Gln Arg Arg 1805 1810 1815

Ser Ser Val Lys Arg Thr Arg Tyr Tyr Glu Thr Tyr Ile Arg Ser 1820 1825 1830

Asp Ser Gly Asp Glu Gln Leu Pro Thr Ile Cys Arg Glu Asp Pro 10 1835 1840 1845

Glu Ile His Gly Tyr Phe Arg Asp Pro His Cys Leu Gly Glu Gln 1850 1855 1860

Glu Tyr Phe Ser Ser Glu Glu Cys Tyr Glu Asp Asp Ser Ser Pro 1865 1870 1875

15 Thr Trp Ser Arg Gln Asn Tyr Gly Tyr Tyr Ser Arg Tyr Pro Gly 1880 1885 1890

Arg Asn Ile Asp Ser Glu Arg Pro Arg Gly Tyr His His Pro Gln 1895 1900 1905

Gly Phe Leu Glu Asp Asp Asp Ser Pro Val Cys Tyr Asp Ser Arg
1910 1915 1920

Arg Ser Pro Arg Arg Leu Leu Pro Pro Thr Pro Ala Ser His 1925 1930 1935

Arg Arg Ser Ser Phe Asn Phe Glu Cys Leu Arg Arg Gln Ser Ser 1940 1945 1950

25 Gln Glu Glu Val Pro Ser Ser Pro Ile Phe Pro His Arg Thr Ala 1955 1960 1965

Leu Pro Leu His Leu Met Gln Gln Gln Ile Met Ala Val Ala Gly
1970 1980

Leu Asp Ser Ser Lys Ala Gln Lys Tyr Ser Pro Ser His Ser Thr 30 1985 1990 1995

Arg Ser Trp Ala Thr Pro Pro Ala Thr Pro Pro Tyr Arg Asp Trp 2000 2005 2010

Thr Pro Cys Tyr Thr Pro Leu Ile Gln Val Glu Gln Ser Glu Ala 2015 2020 2025

5 Leu Asp Gln Val Asn Gly Ser Leu Pro Ser Leu His Arg Ser Ser 2030 2035 2040

Trp Tyr Thr Asp Glu Pro Asp Ile Ser Tyr Arg Thr Phe Thr Pro 2045 2050 2055

Ala Ser Leu Thr Val Pro Ser Ser Phe Arg Asn Lys Asn Ser Asp 10 2060 2065 2070

Lys Gln Arg Ser Ala Asp Ser Leu Val Glu Ala Val Leu Ile Ser 2075 2080 2085

Glu Gly Leu Gly Arg Tyr Ala Arg Asp Pro Lys Phe Val Ser Ala 2090 2095 2100

15 Thr Lys His Glu Ile Ala Asp Ala Cys Asp Leu Thr Ile Asp Glu 2105 2110 2115

Met Glu Ser Ala Ala Ser Thr Leu Leu Asn Gly Asn Val Arg Pro 2120 2125 2130

Arg Ala Asn Gly Asp Val Gly Pro Leu Ser His Arg Gln Asp Tyr 20 2135 2140 2145

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15	Pro			ctg Leu													333
				ctg Leu 40						_			_				381
20				atc Ile													429
				ctg Leu				_	_	_				-		_	477
25				aag Lys													525
30				aag Lys													573
				agc Ser 120													621
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				gcc Ala 200													861
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	_	-		_			_				_			gcc Ala 450	_		1581
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			_	_							_	_	_	aag Lys	_		1677
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15	Cys													gtg Val			1773
		_	_	_	_	_					_	_	_	atc Ile 530	_		1821
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														gtg Val 610			2061
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		gcc Ala															2301
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	-	ccg Pro		_	_	_		_	_				_	_			2541
20		ccg Pro															2589
		gtc Val 790															2637
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		gcc Ala															2781
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			_		_			_	_		_			aag atq Lys Me	-	3117
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				acg Thr 1255							_			cag Gln 1265	gcc Ala	4023
25				gtg Val 1270											tac Tyr	4068
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			_	atc Ile 1300		_					_			_	cag Gln	4158
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				ctc Leu 1375											tac Tyr	4383
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5 Val Lys Ala Leu Arg Ala Phe Arg Val Leu Arg Pro Leu Arg Leu Val

Ser Gly Val Pro Ser Leu Gln Val Val Leu Asn Ser Ile Phe Lys Ala 180 185 190

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Thr Cys Tyr Tyr Ile Gly Thr Asp Ile Val Ala Thr Val Glu Asn Glu 225 230 235 240

15 Lys Pro Ser Pro Cys Ala Arg Thr Gly Ser Gly Arg Pro Cys Thr Ile
245 250 255

Asn Gly Ser Glu Cys Arg Gly Gly Trp Pro Gly Pro Asn His Gly Ile
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Thr His Phe Asp Asn Phe Gly Phe Ser Met Leu Thr Val Tyr Gln Cys 20 275 280 285

Ile Thr Met Glu Gly Trp Thr Asp Val Leu Tyr Trp Val Asn Asp Ala 290 295 300

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25 Gly Ser Phe Phe Ile Leu Asn Leu Val Leu Gly Val Leu Ser Gly Glu 325 330 335

Phe Thr Lys Glu Arg Glu Lys Ala Lys Ser Arg Gly Thr Phe Gln Lys 340 345 350

Leu Arg Glu Lys Gln Gln Leu Glu Glu Asp Leu Arg Gly Tyr Met Ser 30 365 365

- Trp Ile Thr Gln Gly Glu Val Met Asp Val Glu Asp Leu Arg Glu Gly 370 375 380
- Lys Leu Ser Leu Glu Glu Gly Gly Ser Asp Thr Glu Ser Leu Tyr Glu 385 390 395 400
- 5 Ile Glu Gly Leu Asn Lys Ile Ile Gln Phe Ile Arg His Trp Arg Gln 405 410 415
 - Trp Asn Arg Val Phe Arg Trp Lys Cys His Asp Leu Val Lys Ser Arg 420 425 430
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 - Ile Ala Ser Glu His His Asn Gln Pro Leu Trp Leu Thr His Leu Gln 450 455 460
 - Asp Ile Ala Asn Arg Val Leu Leu Ser Leu Phe Thr Ile Glu Met Leu 465 470 475 480
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Gln Val Leu Thr Gly Glu Asp Trp Asn Ser Val Met Tyr Asn Gly Ile 610 615 620

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Ala Gln Lys Ala Lys Ala Glu Glu Arg Lys Arg Arg Lys Met Ser Arg 10 675 680 685

Gly Leu Pro Asp Lys Thr Glu Glu Lys Ser Val Met Ala Lys Lys 690 695 700

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Lys Ala Val Pro Ile Pro Glu Ala Ser Ser Phe Phe Ile Phe Ser Pro 770 780

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785 790 795 800

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Ala Ala Glu Asp Pro Ile Arg Ala Glu Ser Val Arg Asn Gln Ile Leu 820 825 830

Gly Tyr Phe Asp Ile Ala Phe Thr Ser Val Phe Thr Val Glu Ile Val 30 835 840 845

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- 15 Ser Lys Met Thr Glu Glu Glu Cys Arg Gly Tyr Tyr Tyr Val Tyr Lys
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 - Phe Phe Met Met Asn Ile Phe Val Gly Phe Val Ile Val Thr Phe 1055 1060 1065
- 30° Glu Glu Glu Glu Thr Glu Tyr Lys Asn Cys Glu Leu Asp Lys 1070° 1075 1080°

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Thr Phe Leu Ala Ser Ser Gly Gly Leu Tyr Cys Leu Gly Gly 1205 1210 1215

Cys Gly Asn Val Asp Pro Asp Glu Ser Ala Arg Ile Ser Ser Ala 20 1220 1225 1230

Phe Phe Arg Leu Phe Arg Val Met Arg Leu Ile Lys Leu Ser 1235 1240 1245

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Leu Val Asp Gly Thr Gln Ile Asn Arg Asn Asn Asn Phe Gln Thr $30 \ 1295 \ 1300 \ 1305$

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Thr Asn Phe Ala Tyr Tyr Tyr Phe Ile Ser Phe Tyr Met Leu Cys 1355 1360 1365

Ala Phe Leu Ile Ile Asn Leu Phe Val Ala Val Ile Met Asp Asn 10 1370 1375 1380

Phe Asp Tyr Leu Thr Arg Asp Trp Ser Ile Leu Gly Pro His His 1385 1390 1395

Leu Asp Glu Phe Lys Ala Ile Trp Ala Glu Tyr Asp Pro Glu Ala 1400 1405 1410

15 Lys Gly Arg Ile Lys His Leu Asp Val Val Thr Leu Leu Arg Arg 1415 1420 1425

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25 Glu Leu Arg Ala Ile Ile Lys Lys Ile Trp Lys Arg Thr Ser Met 1490 1495 1500

Lys Leu Leu Asp Gln Val Ile Pro Pro Ile Gly Asp Asp Glu Val 1505 1510 1515

Thr Val Gly Lys Phe Tyr Ala Thr Phe Leu Ile Gln Glu His Phe 30 1520 1525 1530

Arg Lys Phe Met Lys Arg Gln Glu Glu Tyr Tyr Gly Tyr Arg Pro 1535 1540 1545

Lys Lys Asp Thr Val Gln Ile Gln Ala Gly Leu Arg Thr Ile Glu 1550 1560

5 Glu Glu Ala Ala Pro Glu Ile Arg Arg Thr Ile Ser Gly Asp Leu 1565 1570 1575

Thr Ala Glu Glu Leu Glu Arg Ala Met Val Glu Ala Ala Met 1580 1585 1590

Glu Glu Arg Ile Phe Arg Arg Thr Gly Gly Leu Phe Gly Gln Val $10 \ 1595 \ 1600 \ 1605$

Asp Thr $\,$ Phe Leu Glu Arg Thr $\,$ Asn Ser Leu Pro $\,$ Pro $\,$ Val Met Ala $\,$ 1610 $\,$ 1615 $\,$ 1620

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Ala Tyr Gly Asn Ser Asn His Ser Asn Asn Gln Met Phe Ser Ser 20 1670 1675 1680

Val His Cys Glu Arg Glu Phe Pro Gly Glu Ala Glu Thr Pro Ala 1685 1690 1695

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                                              1800
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                          1810
                                              1815
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